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RESULT 1
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AR091102 28 Sequence 1222 from patent US AR091102 AR091102.1 GI:10017857

bp DNA 5994076.

linear

PAT 07-SEP-2000

REFERENCE AUTHORS TITLE JOURNAL FEATURES

Unknown.

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Unclassified.

CE 1 (bases 1 to 28)

ARS Chenchik, A., Jokhadze, G. and Bibilashvilli, R.

Methods of assaying differential expression

E Methods of assaying differential expression

LE Methods of assaying differential expression

ANAL Patent: US 5994076-A 1222 30-NOV-1999;

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SUMMARIES

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Chenchik, A., Jokhadze, G. and Bibilashvilli, R.
Methods of assaying differential expression
Patent: US 5994076-A 1221 30-NOV-1999;
Location/Qualifiers
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Unclassified.

1 (bases 1 to 28)

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Chenchik, A., Jokhadze, G. and Bibilashvilli, R.

Methods of assaying differential expression

Patent: US 635829-A 1222 05-MAR-2002;
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Sequence 1221 from patent
AR091101
AR091101.1 GI:10017856
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l (bases 1 to 26)
Chenchik,A., Jokhadze,G. and Bibilashvilli,R. Methods of assaying differential expression Patent: US 6352829-A 1221 05-MAR-2002;
Location/Qualifiers
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Sequence 122
AR198136
AR198136.1
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1 (bases 1 to 22)

1 (bases 1 to 22)

Ackermann, E.J., Bennett, C.Frank., Dean, N.M. a Antisense modulation of novel anti-apoptotic Patent: US 6001992-A 17 14-DEC-1999;
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Sequence
AR095064
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Unknown.
1 (bases 1 to 20)
Ackermann,E.J., Bennett,C.Frank., Dean,N.M. a
Antisense modulation of novel anti-apoptotic
Patent: US 6001992-A 2 14-DEC-1999;
Location/Qualifiers
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RESULT 9 AR095052/c	Qy 39 GGTGAGCCAGCTCAAGACTT 58	Query Match 2.6%; Score 20; DB 6; Length 20; Best Local Similarity 100.0%; Pred. No. 83; Matches 20; Conservative 0; Mismatches 0; Indels 0; Gaps 0;	source 1.20 forganism="unknown" BASE COUNT 4 a 6 c 5 g 5 t ORIGIN	Unclassified 1 (bases 1 the Ackermann, E) Antisense mode Patent: US 60	AR095051/c LOCUS AR095051 20 bp DNA linear PAT 08-SEP-2000 DEFINITION Sequence 4 from patent US 6001992. ACCESSION AR095051 VERSION AR095051.1 GI:10022553 KEYWORDS Unknown. ORGANISM Unknown.	Qy 19 CATTGCCTCAACAGCTTCAA 38	Query Match 2.6%; Score 20; DB 6; Length 20; Best Local Similarity 100.0%; Pred. No. 83; Matches 20; Conservative 0; Mismatches 0; Indels 0; Gaps 0;	EASE COUNT 5 a 2 c 7 g 6 t ORIGIN	Unclassif E 1 (bases S Ackermann Antisense L Patent: U	Κ	H H U	OY 1 GAGTGAGCATTCTCAGCACA 20 . Db 20 GAGTGAGCATTCTCAGCACA 1 RESULT 7	Query Match 2.6%; Score 20; DB 6; Length 20; Best Local Similarity 100.0%; Pred. No. 83; Matches 20; Conservative 0; Mismatches 0; Indels 0; Gaps 0;	/organism="unknown" BASE COUNT 4 a 5 c 5 g 6 t ORIGIN
FEATURES source	REFERENCE 1 AUTHORS A TITLE A JOURNAL P	S	AR095054/c LOCUS LOCUS DEFINITION ACCESSION VERSION	Qy 91 GAA Db 20 GAA RESULT 11	BASE COUNT ORIGIN Query Match Best Local Matches 2	REFERENCE 1 AUTHORS A TITLE A JOURNAL P FEATURES SOURCE	KEYWORDS SOURCE U	ARCESSION A		Query Match Best Local Matches 2	BASE COUNT ORIGIN	REFERENCE 1 AUTHORS A TITLE A JOURNAL P FEATURES SOURCE	ISM	LOCUS A DEFINITION S ACCESSION A VERSION A

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Unclassified.

1 (bases 1 to 20)

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Ackermann.E.J., Bennett, C.Frank., Dean, N.M. and Marcusson, E.G. Ackermann emodulation of novel anti-apoptotic bcl-2-related propatent: US 6001992-A 7 14-DEC-1999;

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Ackermann,E.J., Bennett,C.Frank., Dean,N.M. and Marcusson,E.G. Antisense modulation of novel anti-apoptotic bcl-2-related proteins Patent: US 6001992-A 5 14-DEC-1999;
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Ackermann,E.J., Bennett,C.Frank., Dean,N.M. and Marcusson,E.G. Antisense modulation of novel anti-apoptotic bcl-2-related propatent: US 6001992-A 6 14-DEC-1999;
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1 (bases 1 to 20)

1 (bases 1 to 20)

Ackermann.E.J., Bennett, C.Frank., Dean, N.M. and Marcusson, E.G.

Ackermann E.J., Bennett, C.Frank., Dean, N.M. and Marcusson, E.G.

Antisense modulation of novel anti-apoptotic bcl-2-related proteins

Patent: US 6001992-A 9 14-DEC-1999;

Patent: US 6001992-A 9 14-DEC-1999;

Patent: US 6001992-A 9 14-DEC-1999;
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Ackermann,E.J., Bennett,C.Frank., Dean,N.M. and Marcusson,E.G. Antisense modulation of novel anti-apoptotic bcl-2-related proteins Patent: US 6001992-A 8 14-DEC-1999;
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Search completed: March 14, 2003, 07:13:17 Job time: 1363.8 secs
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AR095058
AR095058.1 GI:10022567
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Ackermann,E.J., Bennett,C.Frank., Dean,N.M. and Marcusson,E.G.
Antisense modulation of novel anti-apoptotic bcl-2-related proteins
Patent: US 6001992-A 11 14-DEC-1999;
Location/Qualifiers
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Ackermann,E.J., Bennett,C.Frank., Dean,N.M. and Marcusson,E.G.
Antisense modulation of novel anti-apoptotic bcl-2-related proteins
Patent: US 6001992-A 10 14-DEC-1999;
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1: /cgm2_6/ptodata/1,
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3: /cgm2_6/ptodata/1,
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Query Match Best Local Similarity

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ATION: E. 37,620 E. 09096/0020 MBER: 09096/0020 ORMATION: -5070 NO: 1222: NO: 1222: ITS: ITS: Itrs Itrs Itrs Itrs Itrs Itrs Itrs Itrs	GENERAL INFORMATION: APPLICANT: Chenchik, Alex APPLICANT: JOKhadze, George APPLICANT: JOKhadze, George APPLICANT: BIBLIASHVILLI, Robert TITLE OF INVENTION: EXPRESSION NUMBER OF SEQUENCES: 1375 CORRESPONDENCE ADDRESS: ADDRESSEE: Fish & Richardson, P.C. STRRET: 2200 Sand Hill Road, Suite 11 CITY: Menlo Park STATE: CA COUNTRY: US ZIP: 94025 ZIP: 94025 COMPUTER READABLE FORM: MEDIUM TYPE: Diskette COMPUTER: IBM Compatible OPERATING SYSTEM: Windows 5 SOTTWARE: PastSEO for Windows Versio SOTTWARE: PastSEO for Windows Versio APPLICATION NUMBER: US/08/859,998 FILLING DATE: 21-MAY-1997 CLASSIFICATION DATA: APPLICATION DATA: APPLICATION DATA: APPLICATION DATA: APPLICATION DATA: APPLICATION UMBER:	14 1 US-08-598 14 1 US-08-598 14 1 US-08-598 19 US-08-988-117 20 1 US-08-598 20 1 US-08-607 20 2 US-08-607 20 2 US-09-08-825 20 3 US-08-825 20 3 US-08-98-25 20 4 US-09-255 21 4 US-09-25 21 US-08-08-060 22 1 US-08-08-060 22 1 US-08-08-060 23 1 US-08-080 24 US-09-280 25 1 US-08-080 26 1 US-08-080	14 1.8 31 1 US-08-309-
)01	□· Z	-/1 Sequence 1, Appl -48 Sequence 48, Appl -254 Sequence 56, Appl -56 Sequence 60, Appl -60 Sequence 31, Appl -10 Sequence 61, Appl -11 Sequence 61, Appl -12 Sequence 11, Appl -13 Sequence 11, Appl -14 Sequence 21, Appl -15 Sequence 21, Appl -16 Sequence 21, Appl -17 Sequence 21, Appl -18 Sequence 21, Appl -19 Sequence 21, Appl -19 Sequence 31, Appl -19 Sequence 32, Appl -19 Sequence 34, Appl -19 Sequence 34, Appl -19 Sequence 54, Appl	560-7 Sequence 7, Sequence 8,

Db Qγ

Matches

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OTHER INFORMATION: oligonucleotide primer; SEQUENCE DESCRIPTION: SEQ ID NO: 1222: US-09-225-928-1222
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                                                                        Sequence 1221, Appearance Patent No. 599407
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Patent No. 6352829
GENERAL INFORMATION:
APPLICANT: Chenchik, Alex
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Best Local :
   GENERAL INFORMATION:
APPLICANT: Chench:
APPLICANT: Jokhadz
APPLICANT: Bibilas
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INFORMATION FOR SEQ ID NO: 1222:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                              265 TCCGTAGACACTGCCAGAACACTATTCA 292
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME: Field, Bret E.
REGISTRATION NUMBER: 37,620
REFERENCE/DOCKET NUMBER: 099
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-322-5070
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: 08/859,998 FILING DATE: 21-MAY-1997 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/225,928
FILLING DATE: 05-Jan-1999
CLASSIFICATION: <Unknown>
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COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson, P.C.
STREET: 2200 Sand Hill Road, Suite 100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Bibilashvilli, Robert TITLE OF INVENTION: METHOD OF ASSAYING DIFFERENTIAL
                                                                                                                                                                                                                                                                                                                                                                                FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                               MOLECULE TYPE: DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows95
SOFTWARE: FastSEQ for Windows Version
                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: nucleic acid
STRANDEDNESS: single
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STATE: CA
Chenchik, Alex
Jokhadze, George
Bibilashvilli, Robert
                                                                                        Application US/08859998
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                                                                                                                                                                                                                                                                    Score 28; DB; Pred. No. 7.
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7.2e-05;
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; OTHER INFORMATION: oligonucleotide primer
US-08-859-998-1221
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US-09-225-928-1221
                                                                                                                                                                                                                                                                                                                     Sequence 1221, Application US/09225928 Patent No. 6352829 GENERAL INFORMATION: APPLICANT: Chenchik, Alex
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match 3.3%;
Best Local Similarity 100.0%;
Matches 26; Conservative
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FILING DATE: 21-WAY-1997
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows95
SOFTWARE: FastSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ZIP: 94025
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson, P.C.
STREET: 2200 Sand Hill Road, Suite 100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION:
TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
LENGTH: 26 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-322-5070
TELEFAX: 415-854-0875
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: field, Bret E.
REGISTRATION NUMBER: 37,620
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TOPOLOGY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CITY: Menlo Park
                 ZIP: 94025
COMPUTER READABLE FORM:
COMPUTER: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows95
SOFTWARE: FastSEQ for Windows Version 2.0
                                                                                                                                                                            NUMBER OF SEQUENCES: 1375
CORRESPONDENCE ADDRESS:
CORRESPONDENCE ADDRESS:
CORRESPONDENCE ADDRESSE: Fish & Richardson,
STREET: 2200 Sand Hill Road, 6
                                                                                                                                                                                                                                                                  Bibilashvilli, Robert
TITLE OF INVENTION: METHOD OF ASSAYING DIFFERENTIAL
                                                                                                                                            CITY: Menlo Park
STATE: CA
                                                                                                         COUNTRY: US
ZIP: 94025
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b; Pred. No. 0.0
0; Mismatches
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Suite
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Matches

RESULT 3

APPLICATION NUMBER: US/09/225,928 FILING DATE: 05-Jan-1999

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RESULT 6
US-09-226-568-2/c
; Sequence 2, Application US/09226568
; Patent No. 6001992
; GENERAL INFORMATION:
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US-09-226-568-17/c
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APPLICANT: Bennett, C. Frank
APPLICANT: Bennett, C. Frank
APPLICANT: Bennett, C. Frank
APPLICANT: Dean, Nicholas M.
APPLICANT: Marcusson, Eric G.
TITLE OF INVENTION: Antisense Modulation of No. 6001992el Anti-apoptotic
TITLE OF INVENTION: bcl-2-Related Proteins
FILE REFERENCE: ISPH-0337
CURRENT APPLICATION NUMBER: US/09/226,568
CURRENT APPLICATION NUMBER: US/09/226,568
CURRENT FILING DATE: 1999-01-07
NUMBER OF SEQ ID NOS: 39
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 17
LENGTH: 22
TYPE: DNA
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Best Local
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INFORMATION FOR SEQ ID NO: 1221:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                              OTHER INFORMATION: Description of Artificial Sequence: antisense OTHER INFORMATION: sequence
                                                                                                                                                                                                                                                                                                                                                       ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                       FEATURE:
                                                                                                                                                                     663 ACGATTGCCAACACATACTTCT 684
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                                                                                                                               Local
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REGISTRATION NUMBER: 37,620
REFERENCE/DOCKET NUMBER: 09096/002001
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-322-5070
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: 08/859,998 FILING DATE: 21-MAY-1997 ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               l Similarity 100.0%;
26; Conservative (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LENGTH: 26 base pairs TYPE: nucleic acid STRANDEDNESS: single TOPOLOGY: linear
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                                                                                                                                                                                                         ilarity 100.0%;
Conservative
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[00.0%; Pred. No.
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Pred. No.
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0.081;
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GENERAL INFORMATION:

APPLICANT: Ackermann, Elizabeth J.

APPLICANT: Bennett, C. Frank

APPLICANT: Dean, Nicholas M.

APPLICANT: Marcusson, Eric G.

TITLE OF INVENTION: Antisense Modulation of No. 6001992el Anti-apoptotic
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US-09-226-568-4/c
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US-09-226-568-3/c
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CURRENT APPLICATION NUMBER: US/09/226,568
CURRENT FILING DATE: 1999-01-07
NUMBER OF SEQ ID NOS: 39
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 2
                                                                                                                                                                                                                                                                               Query Match
Best Local S
Matches 20
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Best Local S
                                                                                                         Sequence 4, Application US/09226568 Patent No. 6001992
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: Antisense Modulation of No. 6001992el Anti-apoptotic TITLE OF INVENTION: bcl-7-Related Proteins
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION NUMBER: US/09/226,568
CURRENT FILING DATE: 1999-01-07
NUMBER OF SEQ ID NOS: 39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Bennett, C. Frank
APPLICANT: Dean, Nicholas M.
APPLICANT: Marcusson, Eric G
                                                                                                                                                                                                                                                                                                                                                                                                             LENGTH: 20
TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Ackermann, Elizabeth J.
                                                                                                                                                                                                                                                                                                                                                                     OTHER INFORMATION: Description of Artificial Sequence: OTHER INFORMATION: sequence
                                                                                                                                                                                                                                                                                                                                                                                                       FEATURE:
                                                                                                                                                                                                             20 GAGTGAGCATTCTCAGCACA 1
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20; Conserv
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0.85;
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Sequence 6, Application US/09226568
Patent No. 6001992
GENERAL INFORMATION:
APPLICANT: Ackermann, Elizabeth J.
APPLICANT: Bennett, C. Frank
APPLICANT: Bennett, C. Frank
APPLICANT: Marcusson, Eric G.
TITLE OF INVENTION: Antisense Modulation of No. 600
TITLE OF INVENTION: Antisense Modulation of No. 600
TITLE OF INVENTION: Antisense Modulation of No. 600
TITLE REFERENCE: ISPH-0337
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION NUMBER: US/09/226,568
CURRENT FILING DATE: 1999-01-07
NUMBER OF SED ID NOS: 39
SOFTWARE: Patentin Ver. 2.0
SEO ID NO 5
LENGTH: 20
TYPE: DNA
ORGANISM: Artificial Sequence
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US-09-226-568-6/c
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US-09-226-568-5/c
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Best Local Similarity
Matches 20; Conserv
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CURRENT APPLICATION NUMBER: US/09/226,568
CURRENT FILING DATE: 1999-01-07
NUMBER OF SEQ ID NOS: 39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: Antisense Modulation of No. 6001992el Anti-apoptotic TITLE OF INVENTION: bcl-2-Related Proteins FILE REFERENCE: ISPH-0337
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Ackermann, Elizabeth J.
APPLICANT: Bennett, C. Frank
APPLICANT: Dean, Nicholas M.
APPLICANT: Marcusson, Eric G.
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TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
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                                                                                                                                                                                                                                                                                                                                                                66 CACCAGGCAGAAGATGACAG 85
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Pred. No.
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                                                                                          6001992el Anti-apoptotic
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Sequence 7, Application US/09226568
Pattent No. 6001992
GENERAL INFORMATION:
APPLICANT: Ackermann, Elizabeth J.
APPLICANT: Marcusson, Eric G.
APPLICANT: Bennett, C. Frank
APPLICANT: Dean, Nicholas M.
APPLICANT: Dean, Nicholas M.
APPLICANT: Marcusson, Eric G.
TITLE OF INVENTION: Antisense Modulation of No. 6001992el Anti-apoptotic
TITLE OF INVENTION: bcl-2-Related Proteins
FILE REFERENCE: ISPH-0337
CURRENT APPLICATION NUMBER: US/09/226,568
CURRENT APPLICATION NUMBER: US/09/226,568
CURRENT FILING DATE: 1999-01-07
NUMBER OF SEQ ID NOS: 39
SOFTWARE: Patentin Ver. 2.0
                               APPLICANT: Ackermann, Elizabeth J.
APPLICANT: Bennett, C. Frank
APPLICANT: Bennett, C. Frank
APPLICANT: Bennett, C. Frank
APPLICANT: Marcusson, Eric G.
TITLE OF INVENTION: Antisense Modulation of No.
TITLE OF INVENTION: Boll-2-related Proteins
FILE REFERENCE: ISPH-0337
CURRENT APPLICATION NUMBER: US/09/226,568
CURRENT FILING DATE: 1999-01-07
NUMBER OF SEQ ID NOS: 39
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 8
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US-09-226-568-8/c
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; OTHER INFORMATION: Description of Artificial Sequence: antisense; OTHER INFORMATION: sequence
US-09-226-568-7
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US-09-226-568-7/c
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; OTHER INFORMATION: Description of Artificial Sequence: antisense
; OTHER INFORMATION: sequence
US-09-226-568-6
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Best Local :
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TYPE: DNA
ORGANISM: Artificial Sequence
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ORGANISM: Artificial Sequence
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Pred. No.
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CURRENT APPLICATION NUMBER: US/09/226,568
CURRENT FILING DATE: 1990-01-07
NUMBER OF SEQ ID NOS: 39
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 10
LENGTH: 20
TYPE: DNA
ORGANISM: Artificial Sequence
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US-09-226-568-9/c
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; OTHER INFORMATION: Description of Artificial Sequence: antisense; OTHER INFORMATION: sequence
US-09-226-568-8
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US-09-226-568-10/c
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Patent No. 6001992
GENERAL INFORMATION:
APPLICANT: Ackermann, Elizabeth J.
APPLICANT: Bennett, C. Frank
APPLICANT: Bennett, C. Frank
APPLICANT: Dean, Nicholas M.
APPLICANT: Dean, Nicholas M.
APPLICANT: INVENTION: Antisense Modulation of No. 6001992el Anti-apoptotic
TITLE OF INVENTION: bcl-2-Related Proteins
FILE REFERENCE: ISPH-0337
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APPLICANT: Ackermann, Elizabeth J.
APPLICANT: Bennett, C. Frank
APPLICANT: Dean, Nicholas M.
APPLICANT: Marcusson, Eric G.
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Best Local Similarity 100.0%; Pred. No.
Matches 20; Conservative 0; Mismatc
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LENGTH: 20
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CURRENT FILING DATE: 1999-01-07
NUMBER OF SEQ ID NOS: 39
SOFTWARE: Patentin Ver. 2.0
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OTHER INFORMATION: Description of Artificial Sequence: antisense OTHER INFORMATION: sequence -09-226-568-10
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US-09-226-568-11/c
Search completed: March 14, 2003, 08:31:49 Job time : 21.863 secs
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                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: Antisense Modulation of No. 6001992el Anti-apoptotic TITLE OF INVENTION: bcl-2-Related Proteins FILE REFERENCE: ISPH-0337 CURRENT APPLICATION NUMBER: US/09/226,568 CURRENT FILING DATE: 1999-01-07 NUMBER OF SEQ ID NOS: 39 SOFTWARE: Patentin Ver. 2.0 Seq ID NO 11
                                                                                                                    Query Match
Best Local Similarity 100.0%; I
Matches 20; Conservative 0;
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                                                                                                                                                                                                                                                            TYPE: DNA ORGANISM: Artificial Sequence
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/ Cgn2_6/ptodata/2/pubpna/US06_PUBCOMB.seq:*

/ Cgn2_6/ptodata/2/pubpna/US06_PUBCOMB.seq:*

/ Cgn2_6/ptodata/2/pubpna/US07_NEW_PUB.seq:*

/ Cgn2_6/ptodata/2/pubpna/US08_NEW_PUB.seq:*

/ Cgn2_6/ptodata/2/pubpna/US08_PUBCOMB.seq:*

/ Cgn2_6/ptodata/2/pubpna/US08_PUBCOMB.seq:*

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                                        0 US-99-854-883-248
0 US-99-752-639-24
0 US-99-984-198-24
0 US-99-981-536-6
0 US-99-951-536-6
0 US-99-951-536-6
0 US-99-989-381-34
0 US-99-899-381-34
0 US-99-899-381-34
0 US-99-894-699-17
10 US-99-894-695-17
9 US-99-894-695-18-35
9 US-99-94-672-69
9 US-99-982-818-69
9 US-99-982-818-69
9 US-99-800-631-81
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               Sequence 6, Appli
Sequence 24, Appl
Sequence 24, Appl
Sequence 6, Appli
Sequence 6, Appli
Sequence 6, Appli
Sequence 34, Appl
Sequence 57, Appl
Sequence 17, Appl
Sequence 17, Appl
Sequence 60, Appl
Sequence 61, Appl
Sequence 81, Appl
Sequence 81, Appl
Sequence 81, Appl
Sequence 44, Appl
Sequence 44, Appl
                                                                                                                                                                                                                                                                                                      Description
RESULT 2
US-09-854-883-248/c
US-09-854-883-248/c
; Sequence 248, Application US/09854883
; Patent No. US/00055479A1
; GENERAL INFORMATION:
    APPLICANT: Lex M. Cowsert
; APPLICANT: Jacqueline Wyatt
; APPLICANT: Susan M. Freier
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1.7	1.7	1.7	1.7	1.7	1.7	1.7	1.7	1.7	1.7	1.7	1.7	1.7	1.7	1.7	1.7	1.7	1.7	1.7	1.7	1.7	1.7	1.7	1.7	1.7	1.7
50	45	42	32	31	31	24	24	23	23	23	23	23	23	23	23	23	23	23	23	23	23	23	23	23	23
9	9	9	10	10	10	10	9	10	10	10	10	10	10	10	10	10	10	10	10	9	9	9	9	9	9
US-09-992-598-192	US-09-747-377-359	US-10-206-654-1	us-09-930-251-9	US-09-782-650-19	-650-	US-09-860-793-16	US-10-167-528-2	US-09-943-851A-20	US-09-944-654-20	· N	-432-2	2	US-09-944-396-20	US-09-945-015-20	587	US-09-944-457-20	US-09-944-449-20	US-09-866-028-20	US-09-841-366A-58	US-10-024-018-1	US-09-944-929-20	US-09-944-907-20	US-09-944-944-20	-944-896	US-09-944-403-20
Sequence 192, App	5	Sequence 1, Appli	Sequence 9, Appli	Sequence 19, Appl	`	•	Sequence 2, Appli	,	•	20,	20,	Sequence 20, Appl	•	Ç		`	•	`	`	Ť	Sequence 20, Appl		Sequence 20, Appl	Sequence 20, Appl	

ALIGNMENTS

; FEATURE: OTHER INFORMATION: Description of Artificial Sequence: Artificial ; OTHER INFORMATION: primer US-09-834-721-6 RESULT 1 US-09-834-721-6/c PRIOR APPLICATION NUMBER: DE 100 26 494.8 PRIOR FILING DATE: 2000-05-27 PRIOR APPLICATION NUMBER: DE 101 02 823.7 PRIOR FILING DATE: 2001-01-23 NUMBER OF SEQ ID NOS: 12 SOFTWARE: Patentin Ver. 2.1 SEQ ID NO 6 Sequence 6, Application US/09834721 Patent No. US20020155551A1 GENERAL INFORMATION: Query Match Best Local : Matches APPLICANT: RIEPING, MECHTHILD TITLE OF INVENTION: PROCESS FOR THE FERMENTATIVE PREPARATION OF L-THREONINE FILE REFERENCE: 21123/280169/MAS CURRENT APPLICATION NUMBER: US/09/834,721 CURRENT FILING DATE: 2001-04-16 TYPE: DNA ORGANISM: Artificial Sequence LENGTH: 20 472 GAATGGATAAGGCA 485 |||||||||||||| 17 GAATGGATAAGGCA 4 Local Similarity 100.0%; Ines 14; Conservative 0; 1.8%; Score 14; DB 9; Log; Pred. No. 1.8e+03; 0; Mismatches 0; Length 20 Indels 0; Gaps 0;

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APPLICANT: ROBERT MCKAY
TITLE OF INVENTION: ANTISENSE MODULATION OF PTP1B EXPRESSION
FILE REFERENCE: 1SPH-0576
CURRENT APPLICATION NUMBER: US/09/854,883
CURRENT FILING DATE: 2001-05-14
PRIOR APPLICATION NUMBER: US 09/629,644
PRIOR APPLICATION NUMBER: US 09/629,644
PRIOR FILING DATE: 2000-07-31
PRIOR APPLICATION NUMBER: US 09/487,368
PRIOR FILING DATE: 2000-01-18
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US-09-752-639-24/c
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Best Local 9
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GENERAL IN
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                                                                                                                                                                                                                                                                                                                                                                                                  OPERATING SYSTEM: WINDOWS
SOFTWARE: FRSTSED for WINDOWS Version 2.0b
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/752,639
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Gatanaga, T.
APPLICANT: Granger, G.A.
TITLE OF INVENTION: Factors Altering Tumor Necrosis
TITLE OF INVENTION: Factor Receptor Releasing Enzyme Activity, and Methods
TITLE OF INVENTION: of Use Thereof
NUMBER OF SEQUENCES: 154
CORRESPONDENCE ADDRESS:
                                   REFERENCE/DOCKET NUMBER: 22
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650-813-5600
                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION DATA: APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM: MEDIUM TYPE: Diskette
                                                                                                                         ATTORNEY/AGENT INFORMATION: NAME: Wu, Frank
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CITY: Palo Alto
STATE: CA
TELEPHONE: 050 - 1792
                                                                                                                                                                  FILING DATE:
APPLICATION NUMBER: 08/964,747
FILING DATE: 05-NOV-1997
APPLICATION NUMBER: 60/030,761
FILING DATE: 06-NOV-1996
                                                                                                      REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                             APPLICATION NUMBER: 09/081,385
                                                                                                                                                                                                                                                                                                                                                                   CLASSIFICATION:
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Madeline M. Butler
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755 PAGE MILL ROAD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   USA
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                                                                                                 41,386
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Pred. No. 1.8e+03;
                                                                                 22000-20577.21
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                                                                      US-09-984-198-24
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US-09-984-198-24/c
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Best Local Similarity 100.0%;
Matches 14; Conservative
Matches 14;
             Query Match 1.8%;
Best Local Similarity 100.0%;
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TELEX: 706141
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS: LENGTH: 20 base pairs
                                                                                                                   SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Granger, G.A.

TITLE OF INVENTION: Factor Receptor Releasing Enzyme Activity, and Methods
TITLE OF INVENTION: of Use Thereof

NUMBER OF SEQUENCES: 154
CORRESPONDENCE ADDRESS:
ADDRESSEE: MORRISON & FOERSTER
STREET: 755 PAGE MILL ROAD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
                                                                                                                                                                                                                        NAME: Wu, Frank
REGISTRATION NUMBER: 41,386
REFERENCE/DOCKET NUMBER: 22000-20577.21
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650-813-5600
                                                                                                                                                                                                                                                                                            FILING DATE: 05-NOV-1997
APPLICATION NUMBER: 60/030,761
FILING DATE: 06-NOV-1996
ATTORNEY/AGENT IMPORMATION:
NAME: Wu, Frank
                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION DATA: APPLICATION NUMBER: 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows
SOSTWARE: FastESQ for Windows Version 2.0b
CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STREET: /--
CITY: Palo Alto
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TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                LENGTH: 20 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                           TELEPHONE: 650-02-
                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: 09/081,385 FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE:
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CLASSIFICATION:
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Conservative
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0; Mismatches
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                 Score 14; DB 10;
Pred. No. 1.8e+03;
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Pred. No. 1.8e+03;
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                                DB 10;
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Gaps
0;
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341 GAATTGTAACCATA 354

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APPLICANT: EEGELING, LOTHAR
APPLICANT: EEGELING, LOTHAR
APPLICANT: SAHM, HERMANN
ITILE OF INVENTION: NEW NUCLEOTIDE SEQUENCES CODING FOR THE THRE GENE
ITILE OF INVENTION: AND PROCESS FOR THE ENZYMATIC PRODUCTION OF
ITILE OF INVENTION: AND PROCESS FOR THE ENZYMATIC PRODUCTION OF
ITILE OF INVENTION: L.THREGNINE USING CORVNEFORM BACTERIA
FILE REFERENCE: 21123/MAS
CURRENT APPLICATION NUMBER: US/09/963,521
CURRENT APPLICATION NUMBER: US/09/963,521
CURRENT FILING DATE: 2001-09-27
PRIOR APPLICATION NUMBER: 09/431,099
PRIOR APPLICATION NUMBER: DE 199 41 478.5
PRIOR FILING DATE: 1999-01
NUMBER OF SEQ ID NOS: 10
SOFTWARE: PATENTIN VET. 2.1
SEQ ID NO 6
LENGTH: 20
TYPE: DNA
AMAGEMENT OF SERVICES OF SERVIC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 5
US-09-951-536-6/c
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US-09-963-521-6/c
                                                                                                                                                     ; OTHER INFORMATION: Description of Artificial Sequence: Primer US-09-963-521-6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 6, Application US/09951536 Patent No. US20020107378A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 6
LENGTH: 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
Query Match 1.8%;
Best Local Similarity 100.0%;
Matches 14; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT:
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                                                                                                                                                                                                                                                                     ORGANISM: Artificial Sequence
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Local Similarity 100.0%; Pred. No. 1.8e+03
nes 14; Conservative 0; Mismatches
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SAHM, HERMANN
THIERBACH, GEORG
            0;
                                         Score 14; DB 10; 1; Pred. No. 1.8e+03;
            Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        red. No. 1.8e+03; Mismatches 0;
                0;
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                                                                                       Length 20
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                Indels
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                0;
                Gaps
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CURRENT APPLICATION NUMBER: US/09/899,381
CURRENT FILING DATE: 2001-07-05
PRIOR APPLICATION NUMBER: 09/398,399
PRIOR FILING DATE: 1999-09-17
NUMBER OF SEQ ID NOS: 53
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 34
LENGTH: 25
TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 7
US-09-398-399-34
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                                QΥ
                                                                                                                                      PEATURE:
OTHER INFORMATION: synthetic probe US-09-899-381-34
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION APPLICANT: DELENSTA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 34, Application Upatent No. US20020068293A1
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: LUEBKE, KEVIN J.
APPLICANT: OVERMAN, LESLIE B.
APPLICANT: SAMPRAS, NICHOLAS M.
APPLICANT: SAMPSON, JEFFREY R.
APPLICANT: WOLBER, PAUL K.
                                                                                     Query Match
Best Local (
                                                                    Matches
                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Wolber, Pual K.
APPLICANT: Sana, Theodore R.
TITLE OF INVENTION: Arrays Having Background Features and
TITLE OF INVENTION: Methods for Using the Same
FILE REFERENCE: 10010760-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: LEFKOWITZ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Delenstarr, Glend C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FEATURE:
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                                   531 ATCTGGCTGGATGA 544
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   ATCTGGCTGGATGA 25
                                                                    14;
                                                                    h Similarity 100.0%; E
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s20020051973A1
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100.0%; Pred. No. 1.8e+0;
ive 0; Mismatches
                                                                                     Score 14; DB 10; 1; Pred. No. 1.8e+03;
                                                                       Mismatches
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RESULT 10
US-09-804-690-17
; Sequence 17, Application US/09804690
; Patent No. US20020034743A1
; GENERAL INFORMATION:
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US-10-083-168-57
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: Endogenous And No. US20030023069A1-Endogenous, Constitutively Act TITLE OF INVENTION: Receptors FILE REFERENCE: AREN-0320 CURRENT ENTITION NUMBER: US/10/083,168 CURRENT ENTLING DATE: 2002-02-26 NUMBER OF SEQ ID NOS: 102 SOFTWARE: Patentin version 3.1 SEQ ID NO 57 LENGTH: 27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION
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Best Local :
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APPLICANT: Chalmers, Derek T.
APPLICANT: Behan, Dominic P.
APPLICANT: Maclejewski-Lenior, Dominique
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: DNA
ORGANISM: Unknown
                                                                                                                                                             COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIAN PC-DOS/MS-DOS
SOFTWARE: PATENTIAN DATA:

CURRENT APPLICATION NUMBER: US/09/804,690
FILING DATE: 12-Mar-2001
CLASSIFICATION: CUNKNOWND-
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/146,187
FILING DATE: CUNKNOWN'
ATTORNEY_AGENT INFORMATION:
NAME: SHERWOOD, Pamela J.
RESISTRATION NUMBER: 36,677
TELECOMMUNICATION INFORMATION:
TELEFHONE: 415-781-1989
TELEFAX: 415-388-3249
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       42 GAGCCAGCTCAAGA 55
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ADDRESSEE: FISH AND RICHARDSON, P.C.
STREET: 2200 SAND HILL ROAD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COHEN, Stanley N
TITLE OF INVENTION: MAMMALIAN TUMOR SUSCEPTIBILITY GENES AND
THEIR USES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: LI, Limin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES: 20
                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                             CITY: MENLO PARK
STATE: CA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Application US/10083168
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Pred. No. 1.9e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 27
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RESULT 11
US-10-092-140-4
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                                                          US-10-092-140-4
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Query Match
Best Local Similarity
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Best Local :
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SEQUENCE CHARACTERISTICS:
LENGTH: 27 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                              TELEX: <Unknown>
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US/09/328,750
FILING DATE: 09-Jun-1999
APPLICATION NUMBER: 60/033,374
FILING DATE: 16-DEC-1996
APPLICATION NUMBER: 60/047,322
FILING DATE: 21-MAY-1997
ATTORNEY/AGENT INFORMATION:
NAME: C"t, France
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              12 GAGCCAGCTCAAGA 25
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                                                                                                                                                                                                                                                    REGISTRATION NUMBER: 4166
REFERENCE/DOCKET NUMBER: 1
TELECOMMUNICATION INFORMATION:
TELEPHONE: 514 845-7126
TELEPAX: 514 288-8389
                                                                           TOPOLOGY: linear
MOLECULE TYPE: Genomic DNA
SEQUENCE DESCRIPTION: SEQ ID NO: 4:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/092,140
FILING DATE: 06-Mar-2002
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ZIP: H3A 2Y3
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
ADDRESSEE: SWABEY OGILVY RENAULT
STREET: 1981 McGill College Avenue - Suite 1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TOPOLOGY: linear
MOLECULE TYPE: other nucleic acid
DESCRIPTION: /desc = "Primer"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: McGill University et al. TITLE OF INVENTION: HUMAN AND MAMMALIAN DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE DESCRIPTION: SEQ ID NO: 17:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES:
                                                                                                                                 LENGTH: 30 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ for Windows Version
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  1.8%;
100.0%;
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  Score 14;
Pred. No.
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. 1.9e+03;
    DB 9; L . 1.9e+03;
                     Length 30;
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Matches

14;

US-09-909-567B-35

Sequence 35, Appropriate Publication No.

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; FEATURE:
; OTHER INFORMATION: Synthetic
US-09-909-567B-35
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APPLICANT: Main, Manoj
APPLICANT: Chen, Seiyu
TITLE OF INVENTION: Compositions and Methods Relating to Lung Specific Genes
FILE REFERENCE: DEX-0214
CURRENT APPLICATION NUMBER: US/09/909,567B
CURRENT APPLICATION NUMBER: 60/219,834
PRIOR FILING DATE: 2001-07-21
PRIOR FILING DATE: 2000-07-21
NUMBER OF SEQ ID NOS: 56
SOFTWARE: Patentin version 3.1
SEQ ID NO 35
LENGTH: 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 60, Applicati
Publication No. US2002
GENERAL INFORMATION:
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                                                       COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: 1BM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION DATA: US/09/734,672
FILING DATE: 03-Dec-2000
    PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 08/966,436
FILING DATE: 07-No. US20020183268A1-97
                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
CORRESPONDENCE ADDRESS:
ADDRESSE: Morgan Lewis & Bockius LLP
STREET: 1111 Pennsylvania Ave., N.W.
                                                                                                                                                                                                                                                                                                                                BRCAL Gene NUMBER OF SEQUENCES: 72
                                                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION: Coding Sequences of the Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Murphy, Patricia D. Allen, Antonette C.
                                                                                                                                                                                                                                CITY: Washington
STATE: District of Columbia
COUNTRY: USA
                                                                                                                                                                                                                ZIP: 20004
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                                                                                                                                                                                                                                                                                                                                                                                                                        Schelter, Denise B.
                                                                                                                                                                                                                                                                                                                                                                                                                                        Critz, Brenda S.
Olson, Sheri J.
                                                                                                                                                                                                                                                                                                                                                                                                     Zeng,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Alvares, Christopher P.
                                                                                                                                                                                                                                                                                                                                                                                                     Bin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 13;
Pred. No.
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. 6.1e+03;
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; SEQUENCE DESCRIPTION: SEQ ID NO: 60:
US-09-734-672-60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 14
US-09-982-828-60/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    REGISTRATION NUMBER: 43,210
REFERENCE/DOCKET NUMBER: 44921
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-739-3000
TELEPAX: 202-739-3001
INFORMATION FOR SEQ ID NO: 60:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
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        PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 09/074,453

FILING DATE: 1998-05-06

APPLICATION NUMBER: US 08/798,691

FILING DATE: 1997-02-12

APPLICATION NUMBER: US 08/598,591

FILING DATE: 1996-02-12

ATTORNEY,AGENT INFORMATION:

NAME: Michael S. Tuscan

REGISTRATION NUMBER: 43,210

REFERENCE/DOCKET NUMBER: 44921-5053-01-US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  494 GCTGGGAAAATGG 506
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nes 13; Conserv
                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
ADDRESSEE: Morgan Lewis & Bockius LLP
STREET: 1111 Pennsylvania Avenue N. W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TOPOLOGY: linear MOLECULE TYPE: DNA (genomic) ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Murphy, Patricia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ATTORNEY/AGENT INFORMATION:
NAME: Michael S. Tusca
                                                                                                                                                                                                                    APPLICATION NUMBER: US/09/982,828
FILING DATE: 22-Oct-2001
CLASSIFICATION: CUnknown>
                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                    STATE: DC
                                                                                                                                                                                                                                                                                                                                                                                                                                              CITY: Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LENGTH: 20 base pairs
TYPE: nucleic acid
STRANDEDNESS: No. US20020183268A1 Relevant
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US 08/598,591 FILING DATE: 12-Feb-96
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INVENTION: Coding Sequences of the Human
BRCA1 Gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Critz, Brenda S. Olson, Sheri J. Thurber, Denise
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Allen, Antonette C.
Alvares, Christopher P.
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Pred. No.
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. 6.2e+03;
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RESULT 13 US-09-734-672-60/c

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US-10-022-819-55/c

Sequence 55, Application US/10022819

Publication No. US20030027166A1

GENERAL INFORMATION:
APPLICANT: ALLEN, Antonette C.
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; SEQUENCE DESCRIPTION: SEQ ID NO: 60:
US-09-982-828-60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              망
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TELEFAX: 202-739-3001
INFORMATION FOR SEQ ID NO: 60:
SEQUENCE CHARACTERISTICS:
                                                                                                                            INFORMATION FOR SEQ ID NO: 55:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                            COMPUTER READBLE FORM:
MEDLUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION WIMBER: US/10/022,819
FILING DATE: 22-Apr-2002
CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
APPLICATION UNMBER: 09/074,452
FILING DATE: 1998-05-06
ATTORNEY/AGENT INFORMATION:
NAME: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          494 GCTGGGAAAATGG 506
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                                         TOPOLOGY: linear
MOLECULE TYPE: other nucleic acid
DESCRIPTION: /desc = "PRIMER"
HYPOTHETICAL: NO
                                                                                                                                                                                                                                          REFERENCE/DOCKET NUMBER: 044921-5049-01-US
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-739-3000
TELEPAX: 202-739-3001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION: CODING SEQUENCE HAPLOTYPE OF THE HUMAN BRCA1 GENE
NUMBER OF SEQUENCES: 67
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS:
ADDRESSEE: Morgan Lewis & Bockius LLP
FRAGMENT TYPE: internal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LENGTH: 20 base pairs
TYPE: nucleic acid
TYPE: nucleic acid
STRANDEDNESS: not relevant
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                                                                                                                                                                                                                                                                                                                                      REGISTRATION NUMBER: <Unknown>
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ANGELLY, Tracy S.
RABIN, Mark B.
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Match
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 ABK67134
ABK67133
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AAT41124
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AAZ39070
AAZ39071
AAZ39072
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Human gene specifi
Human gene specifi
Human Al anti-apop
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Human bcl-xL and b	864(22	20		14	45
Human bcl-xL and b	864	22	20	1.8	14	44
Human bcl-2 mRNA n	540	22	20			43
C. glutamicum tran	124	22	20		14	42
•	2111	22	20		14	41
Newcastle disease	57	21	20		14	40
Homeobox conserved	99	20	20		14	39
RT-PCR primer spec	788	20	20		14	38
PCNA HH ribozyme b	55	22	19		14	37
		22	19		14	36
	ıπ	22	19		14	ω 5
PCNA HH ribozyme b	155	22	19		14	34
PCNA HH ribozyme b		22	19		14	ω w
	639	21	19		14	32
	30	21	19		14	31
HH ribozyme	639	21	19		14	30
	AAA86393	21	19		14	29
PCBA HH ribozyme b	AAA86392	21	19		14	28
Human gene signatu	AAT25577	16	46	•	15	27
Human PTP1b coding	59	22	42		15	26
PTP1B forward PCR	AAZ09965	20	38		15	25
Human PTB1B protei	AAI64146	22	32	•	15	24
Primer 2 relative	ABL40366	24	24	•	15	23
Primer used to amp	4 C	21	20	1.9	15	22
Primer #1 for huma	AAT16171	17	22	•	16	21
Human Al anti-apop	908	21	20	•	20	20
Human Al anti-apop	AAZ39082	21	20	•	20	19
Human Al anti-apop	908	21	20	•	20	18
Human Al anti-apop	3908	21	20	•	20	17
Human Al anti-apop	3907	21	20		20	16
Human Al anti-apop	AAZ39078	21	20	•	20	15
Human Al anti-apop	AAZ39077	21	20		20	14
-apo	907	21	20	•	20	13
Human A1 anti-apop	3907	21	20	•	20	12
Human Al anti-apop	3907	21	20	2.6	20	11
Human Al anti-apop	AAZ39073	21	20	•	20	10

00000000000

ВP

ALIGNMENTS

Human gene specific PCR primer

#1222.

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RESULT 1
ABK67134/c
ID ABK67134;
XX ABK67134;
XX ABK67134;
XX DT O2-JUL-2002 (first entry)
XX ABK67134;
XX Primer; ss; DNA microarray; ox
XX Human gene specific PCR prime
XX Primer; ss; DNA microarray; ox
XX WS6352829-B1.
XX US6352829-B1.
XX D5-MAR-2002.
XX PD 05-MAR-2002.
XX PD 05-JAN-1999; 99US-0225928.
XX PD 05-JAN-1999; 97US-0859998.
XX PA (CLON-) CLONTECH LAB INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Primer; ss; DNA microarray; differential expression analysis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     97US-0859998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                99US-0225928
                                                                                                                                                                                                                                                                                                                                              Bibilashvilli
                                                                                                                                                                                                                                                                                                                                              R;
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producing sub-population of labeled nucleic acids, useful for analysing differences in RNA profiles between several different physiological sources, using set of distinct gene specific primers -

Example 3; SEQ ID No 1222; 11pp; English.

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RESULT 2
ABK67133
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     between several different physiological sources, where the method comprises producing subpopulation of labeled NAs for the different physiological sources, comprising the populations for each physiological source to identify differences in the population, where the comparison concerns the preferably performed by hybridising the labeled NAs for each of the content that the surface of a substrate to produce a hybridisation pattern for each of the sources, and comparing the patterns for each of the sources, where differential gene expression assays are cuilised in differential expression analysis of diseased a normal tissue e.g. neoplastic a normal tissue, or different tissue e.g. neoplastic a normal tissue, or different tissue or subtissue types. The present sequence is a human gene specific PCR of the printed specification, but was obtained in electronic format directly from USPTO at his patent did not form part or property and the printed specification, but was obtained in electronic format directly from USPTO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
                                               Producing sub-population of labeled nucleic acids, useful for analysing differences in RNA profiles between several different physiological sources, using set of distinct gene specific primers -
      Example 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            conditions to enzymatically generate sub-population of NAS, where each gene specific primer has a sequence complementary to a distinct mRNA, and each labeled NA is generated using a single gene specific primer. The method is useful for producing a sub-population of labeled NAS which is useful for analysing the differences in the RNA profiles
                                                                                                                                                                                           WPI; 2002-314699/35.
                                                                                                                                                                                                                                                        Chenchik A,
                                                                                                                                                                                                                                                                                                                     (CLON-) CLONTECH LAB INC
                                                                                                                                                                                                                                                                                                                                                                                  21-MAY-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                            05-JAN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                05-MAR-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US6352829-B1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Primer; ss; DNA microarray; differential expression analysis; human.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human gene specific PCR primer #1221.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           02-JUL-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABK67133;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABK67133 standard; DNA; 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 http.wipo.seqdata.uspto.gov/sequence.html?DocID=6352829B1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Similarity
      SEQ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BP; 6 A; 4 C; 9 G; 9 T; 0
                                                                                                                                                                                                                                                    Jokhadze G, Bibilashvilli R;
                                                                                                                                                                                                                                                                                                                                                                               97us-0859998
                                                                                                                                                                                                                                                                                                                                                                                                                                               99US-0225928
ID No 1221; 11pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    3.6%;
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
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0.00074
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RESULT 3
AAZ39089/c
ID AAZ39089 standard; DNA; 22
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              between several different physiological sources, where the method CC comprises producing subpopulation of labeled NAs for the different CC physiological sources, comprising the population, where the comparison CC source to identify differences in the population, where the comparison CC is preferably performed by hybridising the labeled NAs for each of the CC distinct physiological sources to an array of probe NAs stably CC associated with the surface of a substrate to produce a hybridisation CC pattern for each of the sources, and comparing the patterns for each of the sources, where differential gene expression assays are CC tissue e.g. neoplastic a normal tissue, or different tissue or subtissue types. The present sequence is a human gene specific PCR CC Note: The sequence data for this patent did not form part CC format directly from USPTO at the printed specification, but was obtained in electronic CC format directly from USPTO at the pattern contracts contracts the produce of the printed specification.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
                                 Antisense oligonucleotides which modulate the expression of novel anti-apoptotic bcl-2-related proteins are useful for inducing apoptosis and treating associated diseases e.g. cancer -
   Example 9; Column
                                                                                                             WPI; 2000-061908/05
                                                                                                                                                                                                                      07-JAN-1999;
                                                                                                                                                                                                                                                          07-JAN-1999;
                                                                                                                                                                                                                                                                                             14-DEC-1999
                                                                                                                                                                                                                                                                                                                                US6001992-A
                                                                                                                                                                                                                                                                                                                                                                                     Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                  Human; A1; anti-apoptotic; bcl-2-related protein; antisense inhibition; mcl-1; apoptosis; cancer; antiinflammatory; cytostatic; tumour; inflammation; diagnosis; phosphorothioate; PCR; probe; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human Al anti-apoptotic bcl-2-related protein PCR probe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAZ39089;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention relates to producing a sub-population of labeled nucleic acids (NAs) comprising contacting a NA sample from a physiological source, with a pool of 50 distinct gene specific primers under suitable conditions to enzymatically generate sub-population of NAs, where each gene specific primer has a sequence complementary to a distinct
                                                                                                                                                                                 (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 29-FEB-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           http.wipo.seqdata.uspto.gov/sequence.html?DocID=6352829B1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TCCACCAGGCAGAAGATGACAGACTG
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26; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    nd each labeled NA is generated using a single gene specific The method is useful for producing a sub-population of labeled ch is useful for analysing the differences in the RNA profiles
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ilarity 100.0%;
Conservative 0;
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                                                                                                                                                                                                                                                          99US-0226568
28; 28pp; English
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                                                                                                                                                EG,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ВP
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                                                                                                                                                Bennett CF,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   89
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RESULT 4
AAT41123
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Best Local S
Matches 22
Primers T41001-T41382 are derived from novel human gene signature (GS) sequences which did not match with sequences deposited in Genbank release 76. The GS sequences (T19001-T26837) were obtained from 3'-directed cDNA libraries prepared from various human tissues; synthesis of cDNA was initiated from the 3'-end of mRNA by using poly(T) as the sole primer. Each library is constructed so as to reflect accurately the relative abundance of different mRNAs in the particular tissue from which it was derived. The appearance frequency of a given GS in a cDNA library can be determined (esp. using primers and probes derived from the GS sequences)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present invention describes antisense oligonucleotides which modulate the expression of novel anti-apoptotic bcl-2-related proteins. The antisense oligonucleotides can be used as therapeutic agents to prevent or delay inflammation or tumour formation by promoting apoptosis in human cells or tissues. They can also be used as research agents to establish the function of particular genes and as diagnostic agents in sandwich assays for detecting the level of novel anti-apoptotic bcl-2-related proteins in a sample. The antisense oligonucleotides given in the present invention were designed to target human Al and mcl-1 anti-apoptotic bcl-2-related protein nucleotide sequences. The present sequence represents a PCR probe for the human Al nucleotide
                                                                                                                                                                                                                                                                                                                                            Identifying gene signatures in 3'-directed human cDNA library for diagnosis of abnormal cell function, by preparing cDNA that reflects relative abundance of corresp. mRNA in specific human
                                                                                                                                                                                                                                                                    Example 7; Fig 7; 2245pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matsubara K, Okubo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          12-NOV-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            11-NOV-1994;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            cell typing;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          human;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ignature; messenger RNA; mRNA; relative abundance; cloning; mapping; non-biased library; diagnosis; cyping; abnormal cell function; primer; PCR; amplifi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Similarity 100.
22; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                signature HUMGS01288-derived sense primer.
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0.79;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            amplification;
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                                                                                                                                                                                                                                                                                                                                                                                               e.g
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Вþ δÃ

Query Match Best Local Similarity

2.6%;

Score 20; Pred. No.

DB 16;

Length

20

0

Mismatches

0;

Indels

0;

Gaps

0;

Sequence

20

BP;

ъ P

5 C; 3 G; 7 T; 0 other;

Matches

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8888
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Best Local Si
                    Primers T41001-T41382 are derived from novel human gene signature (GS) sequences which did not match with sequences deposited in Genbank release 76. The GS sequences (T19001-T26837) were obtained from 3'-directed cDNA libraries prepared from various human tissues; synthesis of cDNA was initiated from the 3'-end of mRNA by using poly(T) as the sole primer. Each library is constructed so as to reflect accurately the relative abundance of different mRNAs in the particular tissue from which it was derived. The appearance frequency of a given GS in a cDNA library can be determined (esp. using primers and probes derived from the GS sequences) as a means of diagnosing abnormal cell function or for recognising different cell types. The primers T41123-4 amplify clone pm2245 which comprises the GS HUMGS001288 (T20288), located on chromosome 10.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              as a means of diagnosing abnormal cell function or for recognising different cell types. The primers T41123-4 amplify clone pm2245 which comprises the GS HUMGS001288 (T20288), located on chromosome 10.
                                                                                                                                                                                                                              Example 7; Fig 7;
                                                                                                                                                                                                                                                                        Identifying gene signatures in 3'-directed human cDNA library for diagnosis of abnormal cell function, by preparing cDNA that reflects relative abundance of corresp. mRNA in specific human
                                                                                                                                                                                                                                                                                                                                                                       Matsubara K, Okubo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                01-JUN-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              W09514772-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gene signature; messenger RNA; mRNA; human; cloning; mapping; non-biased
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   03-DEC-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAT41124 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                    12-NOV-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                11-NOV-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        polymerase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      cell typing;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAT41124.;_
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                      (MATS/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           571 TGTGAAATGCTATCTCTÇCT 590
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 TGTGAAATGCTATCTCTCT 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      cloning; mapping; non-biased library;
yping; abnormal cell function; primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Similarity
20; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                    MATSUBARA K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                20 BP; 4 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          chain reaction; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    signature
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                    93JP-0355504.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                94WO-JP01916.
                                                                                                                                                                                                                            2245pp;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    HUMGS01288-derived anti-sense
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                5 C; 3 G;
                                                                                                                                                                                                                              Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (T20288), located
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                8 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    relative abundance; frequency; library; diagnosis; detection; primer; PCR; amplification;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 16;
8.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0,
                                                                                                                                                                                                                                                                                                       e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
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RESULT 7
AAZ39070/C
ID AAZ390
XX
AC AAZ390
XX
DT 29-FEB
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AAZ39069/c
ID AAZ39069 standard; DNA; 20 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Qy
                                                                                                                                                                           Matches
                                                                                                                                                                                                   Query Match
                                                                                                                                                                                                                                                                   The present invention describes antisense oligonucleotides which modulate the expression of novel anti-apoptotic bcl-2-related proteins. The antisense oligonucleotides can be used as therapeutic agents to prevent or delay inflammation or tumour formation by promoting apoptosis in human cells or tissues. They can also be used as research agents to establish the function of particular genes and as diagnostic agents in sandwich assays for detecting the level of novel anti-apoptotic bcl-2-related proteins in a sample. The antisense oligonucleotides given in the present invention were designed to target human Al and mcl-1 anti-apoptotic bcl-2-related protein nucleotide sequences. The present sequence represents an antisense oligonucleotide for the
 29-FEB-2000
                                                    AAZ39070 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Antisense oligonucleotides which modulate the expression of novel anti-apoptotic bc1-2-related proteins are useful for inducing apoptosis and treating associated diseases e.g. cancer - \frac{1}{2}
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human; Al; anti-apoptotic; bcl-2-related protein; antisense inhibition; mcl-1; apoptosis; cancer; antiinflammatory; cytostatic; tumour; inflammation; diagnosis; phosphorothioate; ss.
                           AAZ39070;
                                                                                                                                                                                                                               Sequence 20 BP; 5 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Ackermann EJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  07-JAN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human Al anti-apoptotic bcl-2-related protein antisense oligo #17484.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAZ39069;
                                                                                                                                                                                                                                                          human A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            07-JAN-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     651
                                                                                                                     20
                                                                                                                                                                                          Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ACTGATATGGAAACGATTGC
                                                                                                                                                                           20;
                                                                                                                                                                                          Similarity
                                                                                                                                                                                                                                                         nucleotide sequence.
                                                                                                                                                                           Conservative
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Marcusson EG,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  99US-0226568.
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                                                                                                                                                                                                                                                                                                                                                                                                                                     27;
                                                                                                                                                                                     2.6%;
                                                    DNA; 20
                                                                                                                                                                                                                               2 C; 7 G; 6 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                  28pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         سا
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                                                    ₿P
                                                                                                                                                                          0;
                                                                                                                                                                                       Score 20; DB 21;
Pred. No. 8.1;
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                                                                                                                                                                           Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Dean
                                                                                                                                                                           0;
                                                                                                                                                                                                 Length 20;
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                                                                                                                                                                          Indels
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RESULT 8
AAZ33071,
ID AAZZ3
XX
AC AAZZ
XX
AZ AAZZ
XX
AZ S9-Huma
AX
AX
BE Huma
BE Huma
AX
BE Huma
BE 
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               δÃ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present invention describes antisense oligonucleotides which modulate the expression of novel anti-apoptotic bcl-2-related proteins. The antisense oligonucleotides can be used as therapeutic agents to prevent or delay inflammation or tumour formation by promoting apoptosis in human cells or tissues. They can also be used as research agents to establish the function of particular genes and as diagnostic agents in sandwich assays for detecting the level of novel anti-apoptotic bcl-2-related proteins in a sample. The antisense oligonucleotides given in the present invention were designed to target human Al and mcl-1 anti-apoptotic bcl-2-related protein nucleotide sequences. The present sequence represents an antisense oligonucleotide for the human Al nucleotide sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Antisense oligonucleotides which modulate the expression of novel anti-apoptotic bcl-2-related proteins are useful for inducing apo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human; A1; anti-apoptotic; bcl-2-related protein; antisense inhibition; mcl-1; apoptosis; cancer; antiinflammatory; cytostatic; tumour; inflammation; diagnosis; phosphorothioate; ss.
                                 US6001992-A
                                                                                                                                                                                                                                           Human; A1; anti-apoptotic; bcl-2-related protein; antisense inhibition;
                                                                                                                                                                                                                                                                                                         Human Al anti-apoptotic bcl-2-related protein antisense oligo #17486.
                                                                                                                                                                                                                                                                                                                                                                        29-FEB-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                    AAZ39071;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAZ39071 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 20 BP; 4 A; 6 C; 5 G; 5 T; 0 other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                and treating associated diseases e.g. cancer -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 07-JAN-1999;
                                                                                                                                                                                  inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Ackermann EJ,
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                                                                                      Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (ISIS-) ISIS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                39 GGTGAGCCAGCTCAAGACTT 58
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ò
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GGTGAGCCAGCTCAAGACTT 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PHARM INC
                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                  diagnosis; phosphorothioate; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Marcusson EG,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2.6%; Score 20; DB 21; Length 20; 100.0%; Pred. No. 8.1; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                antiinflammatory;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Bennett CF,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Dean
                                                                                                                                                                                                             cytostatic; tumour;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ¥
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RESULT 9
AAZ390Z
XX AAZ390Z
XX 29-FEE
XX 29-FEE
XX 29-FEE
XX 105-11.
KW Human
KW inflan
XX Synthe
OS Synthe
OS Homo s
XX Synthe
OS 114-DEC
XX ISS
XX 
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Best Local S
Matches 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           the expression of novel anti-apoptotic bcl-2-related proteins. The antisense oligonucleotides can be used as therapeutic agents to prevent or delay inflammation or tumour formation by promoting apoptosis in human cells or tissues. They can also be used as research agents to establish the function of particular genes and as diagnostic agents in sandwich assays for detecting the level of novel anti-apoptotic bcl-2-related proteins in a sample. The antisense oligonucleotides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Antisense oligonucleotides which modulate the expression of novel anti-apoptotic bcl-2-related proteins are useful for inducing apoptosis and treating associated diseases e.g. cancer - \frac{1}{2}
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               given in the present invention were designed to target human A1 and mc1-1 anti-apoptotic bc1-2-related protein nucleotide sequences. The present sequence represents an antisense oligonucleotide for the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAZ39072;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAZ39072 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Ackermann EJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        07-JAN-1999;
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                      WPI; 2000-061908/05
                                                                                                                                              07-JAN-1999;
                                                                                                                                                                                    07-JAN-1999;
                                                                                                                                                                                                                               14-DEC-1999
                                                                                                                                                                                                                                                                     US6001992-A
                                                                                                                                                                                                                                                                                                                                                                                                                    Human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human Al anti-apoptotic bcl-2-related protein antisense oligo #17487.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    29-FEB-2000
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                                                                                                    (ISIS-) ISIS
                                                                                                                                                                                                                                                                                                                                 Synthetic
                                                                                                                                                                                                                                                                                                                                                                         inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           human Al nucleotide sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      present invention describes antisense oligonucleotides which modulate expression of novel anti-apoptotic bcl-2-related proteins. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CACCAGGCAGAAGATGACAG 85
                                                                                                                                                                                                                                                                                                           sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                         Al; anti-apoptotic; bcl-2-related protein; antisense inhibition; apoptosis; cancer; antiinflammatory; cytostatic; tumour;
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20; Conserv
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                                                                                                      PHARM
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                                                              EG,
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Pred.
                                                              Bennett CF,
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No.
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AAZ39073/
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human;
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Antisense oligonucleotides which modulate the expression anti-apoptotic bcl-2-related proteins are useful for indicand treating associated diseases e.g. cancer -
     Column
     27;
28pp; English.
                                                                                sion of novel inducing apoptosis
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mcl-1 anti-apoptotic bcl-2-related protein nucleotide sequences. present sequence represents an antisense oligonucleotide for the establish the function of particular genes and as diagnostic agents sandwich assays for detecting the level of novel anti-apoptotic bcl-2-related proteins in a sample. The antisense oligonucleotides antisense oligonucleotides can be used as therapeutic agents to prevent or delay inflammation or tumour formation by promoting apoptosis in human cells or tissues. They can also be used as research agents to given in the present invention were designed to target human Al and present invention describes antisense oligonucleotides which modulate expression of novel anti-apoptotic bcl-2-related proteins. The A1 nucleotide sequence. in

Sequence 20 BP; 8 A; 4 C; 1 G; 7 T; 0 other;

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Query Match
Best Local :
91 GAATTTGGATATATTTACAG 110
             20;
                    Similarity
              Conservative
                   2.6%;
            0;
                    Score :
              Mismatches
                    . No.
                    DB 2
8.1;
                           21;
              0;
                         Length
              Indels
             0,
              Gaps
              0
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AAZ39073;
                                                                                                              AAZ39073 standard; DNA;
Human Al anti-apoptotic bcl-2-related protein antisense oligo #17488
                                      29-FEB-2000
                                    (first entry)
                                                                                                              20
                                                                                                              ВP
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Synthetic inflammation; apoptosis; A1; anti-apoptotic; bcl-2-related protein; antisense inhibition; diagnosis; phosphorothicate; cancer; antiinflammatory; SS cytostatic; tumour;

sapiens.

US6001992-A.

07-JAN-1999; 99US-0226568

07-JAN-1999; 99US-0226568

(ISIS-) ISIS PHARM INC

Ackermann [] Marcusson EG, Bennett CF, Dean ×

2000-061908/05

Antisense oligonucleotides which modulate the expression of novel anti-apoptotic bcl-2-related proteins are useful for inducing apo and treating associated diseases e.g. cancer apoptosis

Claim 21; Column 27; 28pp; English.

The present invention describes antisense oligonucleotides which the expression of novel anti-apoptotic bcl-2-related proteins. The antisense oligonucleotides can be used as therapeutic agents to go delay inflammation or tumour formation by promoting apoptosis human cells or tissues. They can also be used as research agents establish the function of particular genes and as diagnostic agen o be used as research agents to genes and as diagnostic agents agents to prevent modulate in

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ID AAZ39074 standard;
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       Matches
                    Query Match
Best Local (
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                                                                                                   establish the function of particular genes and as diagnostic agents sandwich assays for detecting the level of novel anti-apoptotic bcl-2-related proteins in a sample. The antisense oligonucleotides given in the present invention were designed to target human Al and mcl-1 anti-apoptotic bcl-2-related protein nucleotide sequences. The present sequence represents an antisense oligonucleotide for the
                                                                                                                                                                                                                                                                                                                                                                      WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            sandwich assays for detecting the level of novel anti-apoptotic bcl-2-related proteins in a sample. The antisense oligonucleotides given in the present invention were designed to target human Al and mcl-1 anti-apoptotic bcl-2-related protein nucleotide sequences. The present sequence represents an antisense oligonucleotide for the human Al nucleotide sequence.
                                                                                                                                                                                                                   The present invention describes antisense oligonucleotides which modulate the expression of novel anti-apoptotic bcl-2-related proteins. The antisense oligonucleotides can be used as therapeutic agents to prevent
                                                                                                                                                                                                                                                                                                          Antisense oligonucleotides which modulate the expression of novel anti-apoptotic bcl-2-related proteins are useful for inducing apoptosis and treating associated diseases e.g. cancer - \frac{1}{2}
                                                              Sequence 20 BP;
                                                                                                                                                                                        or delay inflammation or tumour formation by promoting apoptosis in human cells or tissues. They can also be used as research agents to
                                                                                                                                                                                                                                                                                   Claim 21;
                                                                                                                                                                                                                                                                                                                                                                                                  Ackermann EJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                          07-JAN-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; Al; anti-apoptotic; bcl-2-related protein; antisense inhibition; mcl-1; apoptosis; cancer; antiinflammatory; cytostatic; tumour; inflammation; diagnosis; phosphorothioate; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human Al anti-apoptotic bcl-2-related protein antisense oligo #17489.
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                                                                                                                                                                                                                                                                                                                                                                                                                             (ISIS-) ISIS
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                                                                                          Al nucleotide sequence.
       20;
                       Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Similarity
                                                                                                                                                                                                                                                                                 Column 27;
       Conservative
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Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                  Marcusson EG,
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2.6%; bu
100.0%; Prr
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                                                              4 C; 5 G; 4 T; 0 other;
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Pred. No.
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Pred. No.
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      Mismatches
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                  DB 21
8.1;
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8.1;
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RESULT 12
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ID AAZ390
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Best Local S
Matches 20
AAZ39076
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     or delay inflammation or tumour formation by promoting apoptosis in human cells or tissues. They can also be used as research agents to establish the function of particular genes and as diagnostic agents sandwich assays for detecting the level of novel anti-apoptotic bcl-2-related proteins in a sample. The antisense oligonucleotides given in the present invention were designed to target human Al and mcl-1 anti-apoptotic bcl-2-related protein nucleotide sequences. The
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                                                           AAZ39076 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                present sequence represents an antisense oligonucleotide for the human Al nucleotide sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present invention describes antisense oligonucleotides which modulate the expression of novel anti-apoptotic bcl-2-related proteins. The antisense oligonucleotides can be used as therapeutic agents to prevent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 21; Column
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Antisense oligonucleotides which modulate the expression of novel anti-apoptotic bcl-2-related proteins are useful for inducing apoptosis and treating associated diseases e.g. cancer -
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                                                                                                                                                                                                                                           271 GACACTGCCAGAACACTATT 290
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                                                                                                                                                                                                            GACACTGCCAGAACACTATT 1
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                                                                                                                                                                                                                                                                                                                         . Similarity 20; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                          BP; 4 A; 3 C; 6 G; 7 T; 0 other;
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100.0%; Pi
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                                                           ВÞ
                                                                                                                                                                                                                                                                                                                      Score 20; DB; Pred. No. 8.1 0; Mismatches
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RESULT 14
AAZ39077/c
PN RW KW XXX
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Best Local
                                                                                                                                                                                                                                                                                                                                                      Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present invention describes antisense oligonucleotides which modulate the expression of novel anti-apoptotic bcl-2-related proteins. The antisense oligonucleotides can be used as therapeutic agents to prevent or delay inflammation or tumour formation by promoting apoptosis in human cells or tissues. They can also be used as research agents to establish the function of particular genes and as diagnostic agents in sandwich assays for detecting the level of novel anti-apoptotic cl-2-related proteins in a sample. The antisapes oligonucleotides given in the present invention were designed to target human Al and mcl-1 anti-apoptotic bcl-2-related protein nucleotide sequences. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Antisense oligonucleotides which modulate the expression of novel anti-apoptotic bcl-2-related proteins are useful for inducing apoptosis and treating associated diseases e.g. cancer -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human; Al; anti-apoptotic; bcl-2-related protein; antisense inhibition; mcl-1; apoptosis; cancer; antiinflammatory; cytostatic; tumour; inflammation; diagnosis; phosphorothioate; ss.
                                                                                    Human;
mcl-1;
                                                                                                                                                                                                                                                                                                                                                                                                                                              present s
human Al
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 21; Column 27; 28pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Ackermann
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               07-JAN-1999;
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US6001992-A
                                         Synthetic.
                                                                                                                               Human Al anti-apoptotic bcl-2-related protein antisense oligo #17492.
                                                                                                                                                                                                                     AAZ39077 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                               Sequence
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                           Homo sapiens
                                                                                                                                                            29-FEB-2000
                                                                       inflammation;
                                                                                                                                                                                                                                                                                                           309 GGAGTTTGAAGACGGCATCA 328
                                                                                                                                                                                                                                                                                             20 GGAGTTTGAAGACGGCATCA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Al anti-apoptotic bcl-2-related protein antisense oligo #17491.
                                                                  Al; anti-apoptotic; bcl-2-related protein; antisense inhibition;
apoptosis; cancer; antiinflammatory; cytostatic; tumour;
mation; diagnosis; phosphorothioate; ss.
                                                                                                                                                                                                                                                                                                                                                    Similarity 100.0%; 20; Conservative (
                                                                                                                                                                                                                                                                                                                                                                                                                                                          sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                          sequence represents an antisense oligonucleotide nucleotide sequence.
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Pred. No.
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RESULT 15
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Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    antisense oligonucleotides can be used as therapeutic agents to prevent or delay inflammation or tumour formation by promoting apoptosis in human cells or tissues. They can also be used as research agents to establish the function of particular genes and as diagnostic agents sandwich assays for detecting the level of novel anti-apoptotic bcl-2-related proteins in a sample. The antisense oligonucleotides given in the present invention were designed to target human Al and mol-1 anti-apoptotic bcl-2-related protein nucleotide sequences. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Antisense oligonucleotides which modulate the expression of novel anti-apoptotic bcl-2-related proteins are useful for inducing apoptosis and treating associated diseases e.g. cancer -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI;
                                                                                                                                                                                                                                        Human; A1; anti-apoptotic; bcl-2-related protein; antisense inhibition;
mcl-1; apoptosis; cancer; antiinflammatory; cytostatic; tumour;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               present sequence represents an antisense oligonucleotide for the human {\tt Al} nucleotide sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present invention describes antisense oligonucleotides which modulate the expression of novel anti-apoptotic bcl-2-related proteins. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 21; Column 27; 28pp; English.
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WPI; 2000-061908/05
                                                                             07-JAN-1999;
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                                                                                                                                                                                                  Synthetic
                                                                                                                                                                                                                                                                              Human Al anti-apoptotic bcl-2-related protein antisense oligo #17493.
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                         Ackermann EJ, Marcusson
                                                  (ISIS-) ISIS
                                                                                                       07-JAN-1999;
                                                                                                                                14-DEC-1999.
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                                                                                                                                                                                                                          inflammation; diagnosis; phosphorothioate; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                         379 AAGAAACTTCTACGACAGCA 398
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                                                  PHARM INC
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                           Bennett
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8.1;
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Ax Antisense oligonucleotides which modulate the expression of novel PT anti-apoptotic bcl-2-related proteins are useful for inducing apoptosis PT and treating associated diseases e.g. cancer - XX

Claim 21; Column 27; 28pp; English.

XX

XX

XX

Claim 21; Column 27; 28pp; English.

XX

XX

Che expression of novel anti-apoptotic bcl-2-related proteins. The CC antisense oligonucleotides can be used as therapeutic agents to prevent CC or delay inflammation or tumour formation by promoting apoptosis in CC human cells or tissues. They can also be used as research agents to CC establish the function of particular genes and as diagnostic agents in CC bcl-2-related proteins in a sample. The antisense oligonucleotides CC given in the present invention were designed to target human Al and CC mcl-1 anti-apoptotic bcl-2-related protein nucleotide sequences. The CC present sequence.

XX

Sequence 20 Bp; 5 A; 6 C; 2 G; 7 T; 0 other;

Query Match

Best Local Similarity 100.0%; Pred. No. 8.1;

Matches 20; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

556 GTTACAGGAAAGATCTGTGA 575

Qy 556 GTTACAGGAAAGATCTGTGA 575

Qy 556 GTTACAGGAAAGATCTGTGA 575

Gy GTTACAGGAAAGATCTGTGA 575

Db 20 GTTACAGGAAAGATCTGTGA 14, 2003, 04:46:51

Search completed: March 14, 2003, 04:46:51
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Title:
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OLIGO_NUC Gapop 60.0 , Gapext 60.
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result 0 0 0 0 0 0 0 0 0 0 0 0 ŏ Score Query Match α Length B PCT-US01-47856-2596 PCT-US01-47856-3838 0 US-10-131-827-2596 0 US-10-131-837-3838 0 US-10-131-831-2596 0 US-10-131-831-2596 0 US-10-131-831-2596 0 US-10-131-831-3836 0 US-10-131-831-3836 0 US-09-255-201B-1221 PCT-US99-29593-17 PCT-US99-29593-3 PCT-US99-29593-4 PCT-US99-29593-4 PCT-US99-29593-7 PCT-US99-29593-7 PCT-US99-29593-7 PCT-US99-29593-7 PCT-US99-29593-7 PCT-US99-29593-10 PCT-US99-29593-10 PCT-US99-29593-11 Sequence 2596, Ap Sequence 3838, Ap Sequence 2596, Ap Sequence 2596, Ap Sequence 3838, Ap Sequence 1222, Ap Sequence 1221, Appl Sequence 17, Appl Sequence 17, Appl Sequence 2, Appli Sequence 3, Appli Sequence 4, Appli Sequence 5, Appli Sequence 6, Appli Sequence 7, Appli Sequence 8, Appli Sequence 9, Appli Sequence 9, Appli Sequence 10, Appli Sequence 11, Appli Description

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APPLICANT: Ly, Ngoc
APPLICANT: Ly, Ngoc
APPLICANT: Altman, Peter
ITITLE OF INVENTION: LEUKOCYTE EXPRESSION PROFILING
FILL REFERENCE: 506612000140
CURRENT APPLICATION NUMBER: PCT/US01/47856
CURRENT FILING DATE: 2001-10-22
PRIOR APPLICATION NUMBER: US 60/241,994
PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: US 60/296,764
PRIOR FILING DATE: 2001-6-08
NUMBER OF SEQ ID NOS: 8832
SOFTWARE: Patentin version 3.1
SEQ ID NO 2596
LENGTH: 50
         PCT-US01-47856-3838; Sequence 3838, Application; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                       ; ORGANISM: Homo sapiens PCT-US01-47856-2596
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: BIOCARDIA, INC.
APPLICANT: Wohlgemuth, Jay
APPLICANT: Wohlgemuth, Tay
APPLICANT: Observer, The
APPLICANT: Johnson, Frances
APPLICANT: Fry, Kirk
APPLICANT: Matcuk, George
APPLICANT: Matcuk, George
APPLICANT: Phillips, Julie
APPLICANT: Woodward, Rober
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                                                                                                                                      h 6.4%; Score 50; DB 1; Le Similarity 100.0%; Pred. No. 7.7e-16; 50; Conservative 0; Mismatches 0;
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Phillips, Julie
Woodward, Robert
Ly, Ngoc
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Quertermous, Thomas
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1 PCT-US99-2953-14
1 PCT-US99-2953-16
1 PCT-US99-2953-16
3 US-09-869-894-3
3 US-09-869-894-3
3 US-09-869-894-7
33 US-09-869-894-7
33 US-09-869-894-7
33 US-09-869-894-7
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33 US-09-869-894-11
33 US-09-869-894-12
33 US-09-869-894-12
33 US-09-869-894-13
33 US-09-869-894-15
34 US-09-869-894-15
35 US-09-869-894-15
36 US-09-869-894-16
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38 US-09-869-894-16
39 US-09-869-894-16
30 US-09-869-894-16
30 US-09-869-894-16
31 US-09-869-894-16
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37 US-09-969-894-16
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Sequence 15, Appl
Sequence 16, Appl
Sequence 16, Appl
Sequence 17, Appl
Sequence 17, Appl
Sequence 18, Appl
Sequence 19, Appl
Sequence 19, Appl
Sequence 19, Appl
Sequence 11, Appl
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Sequence 36231, A
Sequence 70874, A
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                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION NUMBER: US 10/006,290
PRIOR APPLICATION NUMBER: US 10/006,290
PRIOR FILLING DATE: 2001-10-22
PRIOR APPLICATION NUMBER: US 60/296,764
PRIOR FILLING DATE: 2001-06-08
NUMBER OF SED ID NOS: 9090
SOFTWARE: Patentin version 3.1
SEQ ID NO 2596
LENGTH: 50
TYPE: DNA
ORGANICAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Ly, NGOC
APPLICANT: Altman, Peter
TITLE OF INVENTION: LEUKOCYTE EXPRESSION PROFILING
FILE REFERENCE: 506612000140
CCURRENT APPLICATION NUMBER: PCT/US01/47856
CCURRENT FILING DATE: 2001-10-22
PRIOR APPLICATION NUMBER: US 60/241,994
PRIOR APPLICATION NUMBER: US 60/241,994
PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: US 60/296,764
PRIOR FILING DATE: 2001-06-08
NUMBER OF SEO ID NOS: 8832
SOFTWARE: Patentin version 3.1
SEQ ID NO 3838
LENGTH: 50
                     RESULT 4
US-10-131-827-3838
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                                                                                                                                                                                                                                                                                          ; ORGANISM: Homo sapiens US-10-131-827-2596
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US-10-131-827-2596
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
APPLICANT: Wohlgemuth, Jay
APPLICANT: FTY, Kirk
APPLICANT: FY, Kirk
APPLICANT: Woodward, Robert
APPLICANT: Ly, Ngoc
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR DIAGNOSING AND MONITORING AUTOIMM
TITLE OF INVENTION: CHRONIC INFLAMMATORY DISEASES
FILE REFERENCE: 506612000120
FILE REFERENCE: 506612000120
Sequence 3838, Application US/10131827
                                                                                                                                                                                                 Query Match 6.4%; Score 50; DB 40; Best Local Similarity 100.0%; Pred. No. 7.7e-16; Matches 50; Conservative 0; Mismatches 0;
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Best Local Similarity 100.0%; Pred. No.
Matches 50; Conservative 0; Mismatcl
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CURRENT FILING DATE: 2002-09-06
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                                                                                                          703 TTGATGATGTAACTTGACCTTCCAGAGTTATGGAAATTTTTGTCCCCATGT 752
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Prentice, James
Phillips, Julie
Woodward, Robert
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Quertermous, Thomas
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PCT-US01-47856-2596

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Query Match Best Local (Matches

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SEQ ID NO 3838
LENGTH: 50
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GENERAL INFORMATION:
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Best Local (
                                                                         Sequence 3838, Application US/10131831 GENERAL INFORMATION:
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Best Local
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SEQ ID NO 2596
LENGTH: 50
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PRIOR FILING DATE: 2001-10-22
PRIOR APPLICATION NUMBER: US 60/296,764
PRIOR FILING DATE: 2001-06-08
NUMBER OF SEQ ID NOS: 9090
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: WOOdward, Robert
APPLICANT: LY, NGOC
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR DIAGNOSING AND MONITORING AUTOIMMUNE
TITLE OF INVENTION: CHRONIC INFLAMMATORY DISEASES
FILE REFERENCE: 506612000120
CURRENT APPLICATION NUMBER: US/10/131,827
CURRENT FILING DATE: 2002-09-06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Wohlgemuth, Jay
APPLICANT: Ery, Kirk
APPLICANT: Woodward, Robert
APPLICANT: Woodward, Robert
APPLICANT: Ly, Ngoc
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR DIAGNOSING AND MONITORING
TITLE OF INVENTION: TRANSPLANT REJECTION
                   APPLICANT: Wohlgemuth, Jay
APPLICANT: Fry, Kirk
APPLICANT: Woodward, Robe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION NUMBER: US 10/006,290 PRIOR FILING DATE: 2001-10-22 PRIOR APPLICATION NUMBER: US 60/296,764
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILE REFERENCE: 506612000121
CURRENT APPLICATION NUMBER: US/10/131,831
CURRENT FILING DATE: 2002-08-05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR FILING DATE: 2001-06-08
NUMBER OF SEQ ID NOS: 9190
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  APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Match 6.4%; Score 50; DB 40; Local Similarity 100.0%; Pred. No. 7.7e-16; es 50; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                        Local Similarity es 50; Conserv
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Woodward, Robert
Ly, Ngoc
              Fry, Kirk
Woodward, Robert
                                                                                                                                                                                                                                                                      6.4%; Sc ilarity 100.0%; P Conservative 0;
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FILE REFERENCE: 50612000121

CURRENT APPLICATION NUMBER: US/10/131/831

CURRENT FILING DATE: 2002-08-05

PRIOR APPLICATION NUMBER: US 10/006,290

PRIOR APPLICATION NUMBER: US 10/006,290

PRIOR FILING DATE: 2001-10-22

PRIOR FILING DATE: 2001-06-08

NUMBER OF SEQ ID NOS: 9190

SOFTWARE: Patentin version 3.1

SEQ ID NO 3838
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; OTHER INFORMATION: oligonucleotide primer; SEQUENCE DESCRIPTION: SEQ ID NO: 1222: US-09-225-201B-1222
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 1222, Application US/US/GENERAL INFORMATION:
APPLICANT: Chenchik, Alex
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
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                                                                                                                                                                                    TELEFAX: 415-854-8975
INFORMATION FOR SEQ ID NO: 1222:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            703 TIGATGATGTAACTIGACCTICCAGAGTTATGGAAATTITGTCCCCATGT
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Local Similarity 100.0%; Pred. No.
nes 50; Conservative 0; Mismatch
                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER: IBM COMPATIBLE
COMPUTER: IBM COMPATIBLE
OPERATING SYSTEM: Windows 95
SOFTWARE: FASTSED for Windows Version
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/225,201B
FILING DATE: 05-Jan-1999
CLASSIFICATION: CUNKNOWN>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ZIP: 94025
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson, P.C.
STREET: 2200 Sand Hill Road, Suite 100
                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: US/08/859,998 FILING DATE: 21-MAY-1997 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                         TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-322-5070
                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION: METHOD OF ASSAYING DIFFERENTIAL
                                                                                         MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                                           NAME: Field, Bret E.
REGISTRATION NUMBER: 37,620
REGISTRATION NUMBER: 09096/002001
                                                                                                                           LENGTH: 28 base pairs TYPE: nucleic acid STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STATE: CA
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                                                                                                                TOPOLOGY:
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Bibilashvilli, R
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7.7e-16;
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PCT-US99-29593-17/c 

: Sequence 17, Application PC/TUS9929593 

: GENERAL INFORMATION:
                                                                                                                                     Дb
                                                                                                                                                                                                                                                                 OTHER IMPORMATION: oligonucleotide primer;
SEQUENCE DESCRIPTION: SEQ ID NO: 1221:
US-09-225-2018-1221
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US-09-225-201B-1221
Sequence 1221, Applicate
GENERAL INFORMATION:
                                                                                   RESULT 9
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Best Local S
Matches 28
                                                                                                                                                                                                                               Query Match
                                                                                                                                                                                                   Matches
APPLICANT: Ackermann, Elizabeth J. APPLICANT: Bennett, C. Frank
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                                                                                                                              265 TCCGTAGACACTGCCAGAACACTATTCA 292
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                                                                                                                                                                                                 Local Similarity es 26; Conserv
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nes 28; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                            REFERENCE/DOCKET NUMBER: 09096/002001
TELEPHONE: 415-322-5070
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US/08/859,998 FILING DATE: 21-MAY-1997 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows95
SOFTWARE: FastSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                   FEATURE:
                                                                                                                                                                                                                                                                                                                             MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Chenchik, Alex
                                                                                                                                                                                                                                                                                                                                                          LENGTH: 26 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                             TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEFAX: 415-854-0875
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: Field, Bret E.
REGISTRATION NUMBER: 37,620
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US/09/225,201B FILING DATE: 05-Jan-1999 CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CITY: Menlo Park
STATE: CA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADDRESSEE: Fish & Richardson, P.C. STREET: 2200 Sand Hill Road, Suite 100
                                                                                                                                                                                               Conservative
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100.0%; Pred. No. 0.00089;
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                                                                                                                                                                                                               Score 26;
Pred. No.
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Isis Pharmaceuticals,

Inc.

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PCT-US99-29593-2/c
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OTHER INFORMATION: Description of Artificial Sequence: antisense sequence
PCT-US99-29593-17
                                                                                                                                     RESULT 11
Sequence 2, Application PC/TUS9929593
GENERAL INFORMATION:
APPLICANT: Ackermann, Elizabeth J.
APPLICANT: Bennett, C. Frank
APPLICANT: Dean, Nicholas M.
APPLICANT: Marcusson, Eric G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION NUMBER: US/09/869,894
CURRENT FILING DATE: 2001-07-06
PRIOR APPLICATION NUMBER: 09/226,568
PRIOR FILING DATE: 1999-01-07
NUMBER OF SEQ ID NOS: 40
SOFTWARE: PatentIN Ver. 2.0
SEQ ID NO 17
                                                                                                                                                                                                                                                                       Matches
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CURRENT FILING DATE: 1999-12-14
EARLIER APPLICATION NUMBER: 09/226,568
EARLIER FILING DATE: 1999-01-07.
NUMBER OF SEQ ID NOS: 40
SOFTWARE: PATENTIAN Ver. 2.0
SEQ ID NO 17
LENGTH: 22
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Best Local Similarity 100.0%; F
Matches 22; Conservative 0;
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APPLICANT:
APPLICANT:
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TYPE: DNA
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APPLICANT: Marcusson, Eric G.
APPLICANT: Isis Pharmaceuticals, Inc.
APPLICANT: Isis Pharmaceuticals, Inc.
TITLE OF INVENTION: Antisense Modulation of Novel Anti-apoptotic bcl-2-Related
TITLE OF INVENTION: Proteins
FILE REFERENCE: ISPH-0432
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TYPE: DNA
ORGANISM: Artificial Sequence
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22; Conserv
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Isis Pharmaceuticals,
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Pred. No.
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Pred. No.
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PCT-US99-29593-3/c
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CURRENT FILING DATE: 1999-12-14
EARLIER APPLICATION NUMBER: 09/226,568
EARLIER FILING DATE: 1999-01-07
NUMBER OF SEQ ID NOS: 40
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 3
                                                                                                                                     Sequence 4, Application PC/TUS9929593 GENERAL INFORMATION:
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LENGTH: 20
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Best Local 9
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APPLICANT: Marcusson, Eric G.
APPLICANT: Isis Pharmaceuticals, Inc.
TITLE OF INVENTION: Antisense Modulation
TITLE OF INVENTION: Proteins
                                                                         APPLICANT: Ackermann, Elizabeth J. APPLICANT: Bennett, C. Frank APPLICANT: Dean, Nicholas M.
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CURRENT FILING DATE: 1999-12-14
EARLIER APPLICATION NUMBER: 09/226,568
EARLIER FILING DATE: 1999-01-07
NUMBER OF SEQ ID NOS: 40
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Dean, Nicholas M.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             LENGTH: 20
TYPE: DNA
ORGANISM: Artificial Sequence
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20; Conserv
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100.0%; Pred. No.
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Pred. No.
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                  of Novel Anti-apoptotic bcl-2-Related
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21;
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21;
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CURRENT APPLICATION NUMBER: PCT/US99/29593
CURRENT FILING DATE: 1999-12-14
EARLIER APPLICATION NUMBER: 09/226,568
EARLIER FILING DATE: 1999-01-07
NUMBER OF SEQ ID NOS: 40
SOFTWARE: PatentIN Ver. 2.0
SEQ ID NO 4
LENGTH: 20
TYPE: DNA
ORGANISM: Artificial Sequence
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EARLIER FILING DATE: 1999-01-07
NUMBER OF SEQ ID NOS: 40
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 5
LENCTH: 20
TYPE: DNA
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Best Local S
Matches 20
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Matches
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TITLE OF INVENTION: Antisense Modulation of TITLE OF INVENTION: Proteins FILE REFERENCE: ISPH-0432 CURRENT APPLICATION NUMBER: PCT/US99/29593 CURRENT FILING DATE: 1999-12-14 EARLIER APPLICATION NUMBER: 09/226,568
                                                                                                                  APPLICANT: Bennett, C. Frank
APPLICANT: Dean, Nicholas M.
APPLICANT: Marcusson, Eric G.
APPLICANT: Isis Pharmaceuticals,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION: Antisense Modulation of Novel Anti-apoptotic bcl-2-Related TITLE OF INVENTION: Proteins FILE REFERENCE: ISPH-0432
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APPLICANT: Bennett, C. Frank
APPLICANT: Dean, Nicholas M.
APPLICANT: Marcusson, Eric G.
APPLICANT: Isis Pharmaceuticals,
                                                                                                                                                                            APPLICANT: Ackermann, Elizabeth J. APPLICANT: Bennett, C. Frank
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FEATURE: OTHER INFORMATION: Description of Artificial Sequence: antisense sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               OTHER INFORMATION: Description of Artificial Sequence: antisense sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                          66 CACCAGGCAGAAGATGACAG 85
                                                                                                                                                                                                                                                                                                                                       20 CACCAGGCAGAAGATGACAG
                                                                                                                                                                                                                                                                                                                                                                                                                                     Local
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                                                                                                                                                                                                                                                                                                                                                                                                                  20;
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                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Application PC/TUS9929593
                                                                                                                                                                                                                                                                                                                                                                                                              2.6%; Score 20; ilarity 100.0%; Pred. No. Conservative 0; Mismatc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
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21;
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21;
                                                                                               Novel Anti-apoptotic bcl-2-Related
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Title:
Perfect score:
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Total number of hits satisfying chosen parameters:
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                                                                                                                                                                                                                                                                                                                                             OM nucleic - nucleic search, using sw model
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                                                                                                                 OLIGO_NUC
Gapop 60.0 , Gapext 60.0
                                                                                                                                                                                                                                                          March 14, 2003, 04:47:29; Search time 213.449 Seconds (without alignments) 8302.956 Million cell updates/sec
                                                                                                                                                                                                      US-09-869-894-1
780
                                                                              4666902 seqs, 1136064160 residues
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                                                                                                                                                                                gagtgagcattctcagcaca.....aattgtatgtatttttctct 780
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  7027196
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Minimum DB seq Maximum DB seq length: 0 length: 50

Post-processing: Listing first 45 summaries

Database : Pending_Patents_NA_New:*

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5: /cgn2_6/ptodata/2/pna/US09_NEW_COMB.seq:*

6: /cgn2_6/ptodata/2/pna/US09_NEW_COMB.seq:*

7: /cgn2_6/ptodata/2/pna/US10_NEW_COMB.seq2:*

7: /cgn2_6/ptodata/2/pna/US10_NEW_COMB.seq2:*

9: /cgn2_6/ptodata/2/pna/US10_NEW_COMB.seq2:*

9: /cgn2_6/ptodata/2/pna/US10_NEW_COMB.seq2:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Score 50 20 20 20 20 20 20 20 20 20 20 20 20 20																									
Match Length DB ID Description 6.4 50 8 US-10-325-899-2596 6.4 50 8 US-10-325-899-3838 2.8 25 9 US-60-427-836-53438 2.7 21 8 US-10-310-188-737 2.4 19 8 US-10-310-188-737 2.4 25 9 US-60-427-836-5872 2.3 25 9 US-60-427-836-5958 2.2 25 9 US-60-427-836-5958 2.2 25 9 US-60-427-836-75564 2.1 25 8 US-10-303-778-10504 2.1 25 8 US-10-355-577-14914 2.1 25 8 US-10-355-577-14914 2.1 25 8 US-10-355-577-48741 2.1 25 8 US-10-355-577-48741 2.1 25 8 US-10-355-577-48741 2.1 25 8 US-10-355-577-48741 2.1 25 8 US-10-355-577-317546 2.1 25 8 US-10-355-577-317546 2.1 25 9 US-60-427-808-35565 2.1 25 9 US-60-427-808-35565 2.1 25 9 US-60-427-808-35565 2.1 25 9 US-60-427-808-35565 2.1 25 9 US-60-427-808-35365 2.1 25 9 US-60-427-808-730126 2.1 25 9 US-60-427-836-242560			N	2	20			_	_	ш	14	13	12				80	7	6	ი . 5	c 4	ω	2	1	Result
Length DB ID Description 50 8 US-10-325-899-2596 50 8 US-10-325-899-3838 50 8 US-10-325-899-3838 50 8 US-10-310-188 Sequence 3334,8 51 9 US-60-427-836-5334,38 Sequence 5334,38 51 9 US-60-427-836-6872 52 9 US-60-427-836-59589 Sequence 6873,7 52 9 US-60-427-808-757664 52 9 US-60-427-808-757664 52 9 US-10-355-577-67556 53 9 US-10-355-577-67556 54 US-10-355-577-149144 Sequence 10504,225 55 9 US-10-355-577-317346 Sequence 24036,235 56 US-10-355-577-317346 Sequence 24036,235 57 9 US-10-355-577-487471 Sequence 24747,256 58 US-10-355-577-487471 Sequence 24747,256 59 US-60-427-808-272157 Sequence 48747,256 59 US-60-427-808-272157 Sequence 272157 50 US-60-427-808-35565 Sequence 242560 50 US-60-427-808-730126 Sequence 730126 50 US-60-427-808-730126 Sequence 57305 50 US-60-427-808-730126 Sequence 57305 50 US-60-427-808-730126 Sequence 57305 50 US-60-427-808-730126 Sequence 57305 50 US-60-427-808-730126 Sequence 242560 50 US-60-427-836-542560 Sequence 2345560 50 US-60-427-836-542560 Sequence 2342560	10	16	16	16	16	16	16	16	16	16	16	16	16	16	16	17	17	18	19	19	21	22	50	50	Score
DB ID Description B US-10-325-899-2596 B US-10-325-899-338 B US-10-325-899-338 B US-10-310-188-53875 B US-10-310-188-53875 B US-10-310-188-5737 B US-60-427-836-59598 B US-60-427-836-59598 B US-60-427-836-59598 B US-60-427-836-757664 B US-10-355-577-67556 B US-10-355-577-10504 B US-10-355-577-149144 B US-10-355-577-149144 B US-10-355-577-317546 B US-10-355-577-3175	2.1	2.1	2.1	2.1	2.1	2.1	2.1	2.1	2.1	2.1	2.1	2.1	2.1	2.1	2.1									6.4	
Description US-10-325-899-2596 US-10-325-899-3838 US-60-427-836-533438 US-60-427-836-533438 US-60-427-836-53559 US-60-427-836-59598 US-60-427-836-33698 US-60-427-836-33698 US-60-427-836-348999 US-60-427-836-348999 US-60-427-836-348999 US-60-427-836-342560	25	1 (51	5	5	5	G	5	5	5	5	G	ъ	G	G	4	J	G	u	J	ø	ш	ū	0	0	1
Description -10-325-899-2596 -10-325-899-3838 -10-310-188-533438 -10-310-188-533438 -10-310-188-53375 -10-310-188-53737 -10-310-188-53737 -10-310-188-53737 -10-310-188-59598 -10-310-188-59598 -10-310-188-59598 -10-310-310-35-59598 -10-310-355-577-149144 -10-355-577-149144 -10-355-577-149144 -10-355-577-149144 -10-355-577-149144 -10-355-577-149144 -10-355-577-149144 -10-355-577-149144 -10-355-577-149144 -10-355-577-149144 -10-355-577-149144 -10-355-577-149144 -10-355-577-149144 -10-355-577-149144 -10-355-577-149144 -10-355-577-149144 -10-355-577-149144 -10-355-577-149144 -10-355-577-149144 -10-355-577-149149 -10-35	-			_	_		_	_			-		-		_					-	-		-	-	;
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Sequence 3	Sequence 3	Sequence 1	Sequence 1	Sequence 3	Sequence 3	Sequence 3	Sequence 3	Sequence 9	Sequence 8	Sequence 7	Sequence 7	Sequence 6	Sequence 4	Sequence 3	Sequence 5	Sequence 9	Sequence 2	Sequence 9	Sequence 5	Sequence 5
398418,	333996,	188567,	172458,	3460, Ap	3459, Ap	382, App	381, App	934308,	842893,	740801,	719610,	686435,	461582,	377737,	51855, A	9159, Ap	2912, Ap	92039, A	55291, A	Appli

ALIGNMENTS

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CURRENT APPLICATION NUMBER: US/10/325,899
CURRENT FILING DATE: 2002-12-20
PRIOR APPLICATION NUMBER: US 60/296,764
PRIOR APPLICATION NUMBER: US 10/206,290
PRIOR FILING DATE: 2001-10-22
PRIOR APPLICATION NUMBER: US 10/006,290
PRIOR FILING DATE: 2001-10-22
PRIOR APPLICATION NUMBER: US 10/131,831
PRIOR FILING DATE: 2002-04-24
NUMBER OF SEQ ID NOS: 9966
SOFTWARE: Patentin version 3.1
SEQ ID NO 2596
LENGTH: 50
TYPE: DNA
ORGANISM: Homo sapiens
US-10-325-899-2596
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 1
US-10-325-899-2596
US-10-325-899-2596, Application US/10325899
GENERAL INFORMATION:
                                                                                                                                                                                                   RESULT 2
US-10-325-899-3838
; Sequence 3838, Application US/10325899
; GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match 6.4\%; Score 50; DB 8; Length 50; Best Local Similarity 100.0\%; Pred. No. 1.2e-16; Matches 50; Conservative 0; Mismatches 0; Indels
APPLICANT: Wohlgemuth, Jay
APPLICANT: Wohlgemuth, Kirk
APPLICANT: Ly, Ngoc
APPLICANT: Ly, Ngoc
APPLICANT: Woodward, Robert
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR DIAGNOSING AND MONITORING TRANSPI
TITLE OF INVENTION: REJECTION
FILE REFERENCE: 506612000122
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Wohlgemuth, Jay
APPLICANT: Fry, Kirk
APPLICANT: Ly, Ngoc
APPLICANT: Woodward, Robert
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR DIAGNOSING AND MONITORING TRANSPL
TITLE OF INVENTION: REJECTION
FILE REFERENCE: 506612000122
                                                                                                                                                                                                                                                                                                                                                                                                                                  703 TTGATGATGTAACTTGACCTTCCAGAGTTATGGAAATTTTGTCCCCATGT 752
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CURRENT APPLICATION NUMBER: US/10/325,899
CURRENT FILING DATE: 2002-12-20
PRIOR APPLICATION NUMBER: US 60/296,764
PRIOR FILING DATE: 2001-06-08
PRIOR APPLICATION NUMBER: US 10/006,290
PRIOR FILING DATE: 2001-10-22
PRIOR APPLICATION NUMBER: US 10/131,831
PRIOR APPLICATION NUMBER: US 10/131,831
PRIOR FILING DATE: 2002-04-24
NUMBER OF SEQ ID NOS: 9966
SOPTWARE: Patentin version 3.1
SEQ ID NO 3838
LENGTH: 50
                                                                                                                                      US-10-310-188-73875/c
Sequence 73875, Application US/10310188
Sequence 73875, Application US/10310188
GENERAL INFORMATION:
APPLICANT: RosettaGemonics
TITLE OF INVENTION: BIOINFORMATICALLY DETECTABLE GROUP OF NOVEL VIRAL REGULATORY GENE
TITLE OF INVENTION USES THEREOF
FILE REFERENCE: 47487
CURRENT APPLICATION NUMBER: US/10/310,188
CURRENT FILLING DATE: 2002-12-19
NUMBER OF SEQ ID NOS: 86841
SOFTWARE: Patentin version 3.1
SEQ ID NO 73875
LENGTH: 21
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                                                                              : LENGTH: 21
; TYPE: DNA
; ORGANISM: Homo sapiens
US-10-310-188-73875
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; ORGANISM: Rattus norvegicus
US-60-427-836-533438
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US-60-427-836-533438
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 53438, application US/60427836
GENERAL INFORMATION:
APPLICANT: Xue Mei Zhou
TITLE OF INVENTION: Methods of Genetic Analysis of Rat
FILE REFERENCE: 3527
CURRENT APPLICATION NUMBER: US/60/427,836
CURRENT FILING DATE: 2002-11-20
NUMBER OF SEQ ID NOS: 699466
SOFTWARE: Microarray Probe Sequence Listing Generator V 1.1
SEQ ID NO 533438
LENGTH: 25
Query Match 2.7%;
Best Local Similarity 100.0%;
Matches 21; Conservative
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Best Local :
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les 50; Conserv
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  0;
                   Score 21; DB 8; Pred. No. 0.8;
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    Mismatches
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0.23;
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                                     Length 21,
    Indels
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Sequence 59598, Application US/60427836
GENERAL INFORMATION:
APPLICANT: Xue Mei Zhou
TITLE DE INVENTION: Methods of Genetic Analysis of Rat
FILE REFERENCE: 3527
CURRENT APPLICATION NUMBER: US/60/427,836
CURRENT FILING DATE: 2002-11-20
NUMBER OF SEQ ID NOS: 699466
SOFTWARE: Microarray Probe Sequence Listing Generator V 1.1
SEQ ID NO 59598
LENGTH: 25
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TYPE: DNA
; ORGANISM: Rattus norvegicus
US-60-427-836-6872
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-60-427-836-6872, Application US/60427836; Sequence 6872, Application US/60427836; GENERAL INFORMATION:
GENERAL INFORMATION: Med Zhou
TITLE OF INVENTION: Methods of Genetic Analysis of
FILE REFERENCE: 3527
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US-10-310-188-85737/c
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GENERAL INFORMATION:
APPLICANT: ROSettaGemonics
TITLE OF INVENTION: BIOINFORMATICALLY DETECTABLE GROUP OF NOVEL VIRAL REGULATORY (
TITLE OF INVENTION: USES THEREOF
FILE REFERENCE: 47487
CURRENT APPLICATION NUMBER: US/10/310,188
CURRENT APPLICATION NUMBER: US/10/310,188
CURRENT FILING DATE: 2002-12-19
NUMBER OF SEQ ID NOS: 86841
SOFTWARE: PatentIn version 3.1
SEQ ID NO 85737
LENGTH: 19
TENGTH: 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQ ID NOS: 699466
SOFTWARE: Microarray Probe Sequence Listing Generator V 1.1
SEQ ID NO 8872
LENGTH: 25
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Best Local Similarity
Matches 19; Conserv
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Best Local Similarity 100.0%;
Matches 19; Conservative
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ORGANISM: Rattus norvegicus
                    TYPE: DNA
                                                                                                                                                                                                                                                                                                                                        173 AAACGTCCAGAGTGCTACA 191
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ilarity 100.0%;
Conservative (
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Pred. No.
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US-10-303-778-10504/c
US-10-303-778-10504/c
Sequence 10504, Application US/10303778
GENERAL INFORMATION:
APPLICANT: RosettaGenomics
APPLICANT: ROSETTAGENOMICS
TITLE OF INVENTION: BIOINFORMATICALLY DETECTABLE GROUP OF NOVEL VIRAL
TITLE OF INVENTION: REGULATORY GENES AND USES THEREOF
FILE REFERENCE: 47416
CURRENT APPLICATION NUMBER: US/10/303,778
CURRENT FILING DATE: 2002-11-26
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Sequence 757664, Application US/60427808
GENERAL INFORMATION:
APPLICANT: Xue Mei Zhou
TITLE OF INVENTION: Methods of Genetic Analysis of Mouse
FILE REFERENCE: 3528
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US-60-427-808-360861
Sequence 360861, Application US/60427808
GENERAL INFORMATION:
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Best Local S
Matches 18
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Matches
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CURRENT FILING DATE: 2002-11-20
NUMBER OF SEQ ID NOS: 982914
SOFTWARE: Microarray Probe Sequence Listing Generator V 1.1
SEQ ID NO 757664
LENGTH: 25
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CURRENT FILING DATE: 2002-11-20
NUMBER OF SEQ ID NOS: 982914
SOFTWARE: Microarray Probe Sequence Listing Generator V 1.1
SEQ ID NO 360861
LENGTH: 25
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Best Local S
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TITLE OF INVENTION: Methods of Genetic Analysis of Mouse
FILE REFERENCE: 3528
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ilarity 100.0%;
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100.0%; Pred. No. 35;
tive 0; Mismatches
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Pred. No.
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35;
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; SOFTWARE: Microarray Probe So; SEQ ID NO 149144
; LENGTH: 25
; TYPE: DNA
; ORGANISM: Homo sapien
US-10-355-577-149144
RESULT 13
US-10-355-577-249093
; Sequence 249093, Application US/10355577
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; TYPE: DNA; ORGANISM: Infectious US-10-303-778-10504
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; TYPE: DNA
; ORGANISM: Homo sapien
US-10-355-577-67556
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GENERAL INFORMATION:
APPLICANT: Mittmann, Michael
TIPLE OF INVENTION: Methods of Genetic Analysis
FILE REFERENCE: 3121
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Mittmann, Michael
TITLE OF INVENTION: Methods of Genetic Analysis of Probes: FILE REFERENCE: 3121
CURRENT APPLICATION NUMBER: US/10/355,577
CURRENT FILING DATE: 2003-01-31
NUMBER OF SEQ ID NOS: 997516
SOFTWARE: Microarray Probe Sequence Listing Generator V 1.1
SEQ ID NO 67556
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LENGTH: 24
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Best Local Similarity
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Best Local
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CURRENT FILING DATE: 2003-01-31
NUMBER OF SEQ ID NOS: 997516
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SOFTWARE: PatentIn version 3.1
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100.0%; Pred. No.
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; APPLICANT: Mittmann, Michael
; TITLE OF INVENTION: Methods of Genetic Analysis of Probes: HG-Ul33
; FILE REFERENCE: 3121
; CURRENT APPLICATION NUMBER: US/10/355,577
; CURRENT FILING DATE: 2003-01-31
; NUMBER OF SEQ ID NOS: 997516
; SOFTWARE: Microarray Probe Sequence Listing Generator V 1.1
; SEQ ID NO 249093
; ELNGTH: 25
; TYPE: DNA
; ORGANISM: Homo sapien
US-10-355-577-249093
                                                                                                                                                                                                                                                                                                                                                  US-10-355-577-487471/c
US-10-355-577-487471/c
Sequence 487471, Application US/10355577
GENERAL INFORMATION:
APPLICANT: Mittmann, Michael
FITLE OF INVENTION: Methods of Genetic Analysis of Probes: HG-Ul33
FILE REFERENCE: 3121
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CURRENT FILING DATE: 2003-01-31
VUMBER OF SEQ ID NOS: 997516
SOPTWARE: Microarray Probe Sequence Listing Generator V 1.1
SEQ ID NO 317546
LENGTH: 25
TYPE: DNA
ORGANISM: Homo sapien
US-10-355-577-317546
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US-10-355-577-317546
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GENERAL INFORMATION:

APPLICANT: Mittmann, Michael

TITLE OF INVENTION: Methods of Genetic Analysis of Probes: HG-Ul33;

FILE REFERENCE: 3121
                                                                                                                                                                                                                      CURRENT APPLICATION NUMBER: US/10/355,577
CURRENT FILING DATE: 2003-01-31
NUMBER OF SEQ ID NOS: 997516
SOFTWARE: Microarray Probe Sequence Listing Generator V 1.1
SEQ ID NO 487471
LENGTH: 25
TYPEE- NUS
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                                                                      2.1%;
Local Similarity 100.0%;
hes 16; Conservation
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ACCAGAAAGGACACTC 6
                                                             Score 16; DB 8; Lengum 20, pred. No. 4.3e+02; riamatches 0; Indels
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Search completed: March 14, 2003, 10:49:52 Job time: 215.449 secs

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Title:
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AUTHORS TITLE JOURNAL FEATURES	REFERENCE	ORGANISM	SOURCE	KEYWORDS	VERSION	ACCESSION	DEFINITION	LOCUS	186713/c	RESULT 1
<pre>Reed, J.C. and Sato, T. Interaction of proteins involved in a cell death pathway Patent: US 5702897-A 1 30-DEC-1997; Location/Qualifiers</pre>	Unclassified. 1 (bases 1 to 21)	Unknown.	Unknown.	•	I86713.1 GI:3206431	I86713	Sequence 1 from patent US 5702897.	I86713 21 bp DNA		
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Ackermann, E.J., Bennett, C.Frank., Dean, N.M. and Marcusson, I
Antisense modulation of novel anti-apoptotic bcl-2-related
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Ackermann,E.J., Bennett,C.Frank., Dean,N.M. and Marcusson,E.G. Antisense modulation of novel anti-apoptotic bcl-2-related proteins Patent: US 6001992-A 23 14-DEC-1999;
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Ackermann, E.J., Bennett, C. Frank., Dean, N.M. a Antisense modulation of novel anti-apoptotic Patent: US 6001992-A 22 14-DEC-1999;
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Sequence
AR095071
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Sequence
AR095072
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1 (bases 1 to 20)

Ackermann, E.J., Bennett, C. Frank., Dean, N.M. and Marcusson, E.G. Antisense modulation of novel anti-apoptotic bcl-2-related propatent: US 6001992-A 25 14-DEC-1999;
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20; Conserv
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1 (bases 1 to 20)

Ackermann, E.J., Bennett, C. Frank., Dean, N.M. and Marcusson, E.G. Antisense modulation of novel anti-apoptotic bcl-2-related proteins Patent: US 6001992-A 24 14-DEC-1999;
                                                                          l Similarity
20; Conserv
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Ackermann, E.J., Bennett, C. Frank., Dean, N.M. a
Antisense modulation of novel anti-apoptotic
Patent: US 6001992-A 26 14-DEC-1999;
                                                                                                                                                                                                                                                      Similarity
20; Conserv
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Ackermann,E.J., Bennett,C.Frank., Dean,N.M. and Marcusson,I
Antisense modulation of novel anti-apoptotic bcl-2-related
Patent: US 6001992-A 27 14-DEC-1999;
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                             1 (bases 1 to 20)
Ackermann, E.J., Bennett, C.Frank., Dean, N.M. and Marcusson, I Antisense modulation of novel anti-apoptotic bcl-2-related Patent: US 600192-A 28 14-DEC-1999;
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Ackermann,E.J., Bennett,C.Frank., Dean,N.M. and Marcusson,E.G.
Antisense modulation of novel anti-apoptotic bcl-2-related proteins
Patent: US 6001992-A 30 14-DEC-1999;
Location/Qualifiers
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Ackermann, E.J., Bennett, C.Frank., Dean, N.M. and Marcusson, E.G.
Antisense modulation of novel anti-apoptotic bcl-2-related proteins
Patent: US 6001992-A 29 14-DEC-1999;
Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match 0.5%; Strest Local Similarity 100.0%; Strest Matches 20; Conservative 0;
                                                                                                                                                                                                                                                                          ORGANISM
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AR095079
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AR095078
AR095078.1
                                                                                               Similarity 20; Conserv
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Ackermann, E. J., Bennett, C. Frank., Dean, N.M. and Marcusson, E.G.
Antisense modulation of novel anti-apoptotic bcl-2-related proteins
Patent: US 6001992-A 32 14-DEC-1999;
                                                                                                                                                                                                                                                                                      Unknown.
                                                                                                                                                                                                                                                                                                                                                    AR095079
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Unclassified.

1 (bases 1 to 20)

Ackermann, E.J., Bennett, C.Frank., Dean, N.M. and Marcusson, E.G. Antisense modulation of novel anti-apoptotic bcl-2-related proteins Patent: US 6001992-A 31 14-DEC-1999;
                                                                                                                                                                                                                                                            Unclassified
                                                                                                                                                                                                                                                                           Unknown
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5 c 3 g
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       from patent US 6001992.
                                                                                              Score 20; DB; Pred. No. 39; 0; Mismatches
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3934
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1: \SIDS2\gcgdata\geneseq\geneseqn-embl\Na1980.DAT: *
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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Human mcl-1 anti-a	Human mcl-1 anti-a	Human Mcl-1 gene r	Human Mcl-1 cDNA a	Human Mcl-1 cDNA a	PCR primer P13. S	PCR primer P12. S	Human Mcl-1 gene f	Human Mcl-1 DNA ex	Description		

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N. meningitidis fu	Nei	Triple helix	Sequence of s	Sequence of s	erb B2/neu pi	Murine Srv	PCR primer SryMm	PCR primer used t	Guanosine rich o	Guanosine r	erb B2/neu promo	GnT-IV cDNA contai	Human Mcl-1 cDNA	Human Mcl-1 cDNA	Human Mcl-1	Human myelo	Human myeloid leuk	an	ant	Human mcl-1 anti-a	Human mcl-1 anti-a	anti-	anti-	•	mcl-1 ant	Human mcl-1 anti-a	Human mcl-1 anti-a	Human mcl-1 anti-a	Human mcl-1 anti-a		Human mcl-1 anti-a	Human mcl-1 anti-a	Human mcl-1 anti-a	-1 ant	Human mcl-1 anti-a	

ALIGNMENTS

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Human Mcl-1 DNA extending PCR primer, #105C.
                                (first entry)
                                                                                                 30
                                                                                                 ВP
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RESULT 1

AAD06442/c

ID AAD06442 standard; DNA; 30

XX

AAD06442;

XX

AAD06442;

XX

PT 10-AUG-2001 (first entry)

DT 10-AUG-2001 (first entry)

DT 10-AUG-2001 (first entry)

DT 10-AUG-2001 (first entry)

XX

Human Mcl-1 DNA extending

XX

Human; Mcl-1 gene regulate

KW neuronal cell; tumour cell

KW neuronal rell; tumour cell

KW neuronal arthritis; sepit

KW neurodegenerative disease;

XX

PN w0200136594-A1.

XX

PD 25-MAY-2001.

XX

PD 25-MAY-2001.

XX

PD 25-MAY-2001.

XX

PR 14-JAN-2000; 2000WO-US0096

XX

PR 16-NOV-1999; 99US-01661:

XX

PA (DART-) DARTMOUTH COLLEGE

XX

PA (DART-) DARTMOUTH COLLEGE Human; Mcl-1 gene regulatory element; Mcl-1s/deltaTM variant; neuronal cell; tumour cell; apoptosis; therapy; cancer; psoriasis; diabetic retinopathy; corneal graft neocularisation; neovascular glaucoma; epithelial condition; autoimmune disease; rheumatoid arthritis; systemic lupus erythematosus; neurodegenerative disease; PCR primer; ss. 14-JAN-2000; 2000WO-US00969. (DART-) DARTMOUTH COLLEGE. 99US-0166113 Whyte ĭ

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RESULT 2
AAT0314/c
ID AAT03114/c
XX AAT03114/c
XX O5-JUN
XX MC1-1;
KW Mc1-1;
KW Mc1-1;
KW Getect
XX Synthe
XX Synthe
PN W09528
XX Synthe
PN W09528
XX II-APR
XX II-APR
XX III-APR
XX III-AP
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                         AATO3164-T031366 are primers used for the amplification of the hum McI-1 gene. The McI-1 protein is a BcI-2 related protein, the McI-1 protein was expressed and used to investigate the interactions of McI-1 with other proteins known to be involved in apoptosis (excluding the Bax protein). Proteins detected using this method (excluding the Bax protein).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present invention relates to Mcl-1 gene regulatory elements and the variant Mcl-1s/deltaTM. The anti-apoptotic Mcl-1 protein is encoded by exons 1, 2 and 3. The pro-apoptotic Mcl-1s/DeltaTM variant encoded by exons 1 and 3 is obtained due to alternative mRNA splicing. The Mcl-1 gene regulatory element is useful for modulating the Mcl-1 gene expression in a cell e.g., neuronal cell or tumour cell, such that apoptosis of the cell is induced or cell viability is increased. The Mcl-1 and its regulatory elements are used for treating pathological conditions which include cancer, diabetic retinopathy, corneal graft neovascularisation and neovascular glaucoma, epithelial conditions such as psoriassis, autoimmune diseases like rheumatoid arthritis, systemic lipus erythematosus, and neurodegenerative diseases. The present sequencis a PCR primer #105C, used for the extension reaction of the human
                                                                                                                                                                                                  Example 1; Page 22; 62pp; English.
                                                                                                                                                                                                                                                                 with proteins involved in apoptosis
                                                                                                                                                                                                                                                                                       Detection of novel proteins involved in apoptosis -
                                                                                                                                                                                                                                                                                                                                                         WPI; 1995-373811/48
                                                                                                                                                                                                                                                                                                                                                                                                                 Reed JC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      detection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (LJOL-) LA JOLLA CANCER RES FOUND
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  13-APR-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           12-APR-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                26-OCT-1995
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mcl-1; Bax; apoptosis; cell death; regulation; Bcl-2; novel;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        30
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upstream activators or downstream effectors of a cellular
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3. 0.00072;
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                         this method car
                                                                                                                                                                                                                                                                                          interaction
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                                                                                                                the Mcl-1
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                                                                                             RESULT 3
AAX26538
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                                                       Query Match
Best Local S
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                                                                                                                           template DNA containing nucleotide analogues is amplified in the presence of nucleotide analogues and a substance which lowers the Tm value of double-stranded nucleic acids. Suitable nucleotide analogues are 7-deaza-dGTP, 7-deaza-dATP, dITP and hydroxymethyl-dUTP. Suitable Tm value-lowering substances are formamide, dimethyl sulphoxide and trimethyl-glycine. The methods improve the amplification of DNA. Also, DNA fragments which originated as RNA can be amplified without
                                                                                           Sequence 21 BP; 4 A; 5 C; 7 G; 5 T; 0 other;
                                                                                                                                                                                                                                                                         Simple and accurate method for DNA amplification - uses amplification in the presence of nucleotide analogues together with a compound which lowers the Tm of double-stranded nucleic acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1235 GCAAGTGGCAAGAGGATTATG 1255
                      586
                                                                                                                   purifying
                                                                                                                                                                                                                PCR primers AAX26538-39 were used to exemplify the inv specification describes methods for DNA amplification,
                                                                                                                                                                                                                                                                                                                                                    Hino F,
                                                                                                                                                                                                                                                                                                                                                                         (TAKI ) TAKARA SHUZO CO LTD
                                                                                                                                                                                                                                                                                                                                                                                                   21-OCT-1997;
14-AUG-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DNA amplification; nucleotide analogue; PCR primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PCR primer P12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    27-MAY-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAX26538
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  protein such as Bax which induces apoptosis. If the BC1-2 related protein apoptosis levels are decreased protein binding to and inactivating Bax.
                                                                                                                                                                                                                                                                                                                                                                                                                                      10-AUG-1998;
                                                        Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                21 GCAAGTGGCAAGAGGATTATG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local
              CGGCAGTCGCTGGAGATTATC
                                                                                                                                                                                                                                                  6; Page 31; 36pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Similarity
                                                                                                                                                                                                                                                                                                                                                    Kato I,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  standard;
                                             Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                  RNAs in sample.
                                                                                                                                                                                                                                                                                                                                                                                                  97JP-0305016
97JP-0231885
                                                                                                                                                                                                                                                                                                                                                                                                                                      98WO-JP03566
                                                                                                                                                                                                                                                                                                                                                    Mukai H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DNA;
                                                        100.0%;
                                                                     0.5%;
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                                                       Score 21;
Pred. No.
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Pred. No.
                                             Mismatches
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                                                        DB 20;
. 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               If the protein ecreased due to
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                                                                 Length 21;
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                                            Indels
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                                                                                                                                                                                                                            invention.
                                                                                                                                                                                                                wherein
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                                          Gaps
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RESULT 4

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RESULT 5
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ID AAX26539 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local S
Matches 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PCR primers AAX26538-39 were used to exemplify the invention. The specification describes methods for DNA amplification, wherein a template DNA containing nucleotide analogues is amplified in the presence of nucleotide analogues and a substance which lowers the Tm value of double-stranded nucleic acids. Suitable nucleotide analogues are 7-deaza-dGTP, 7-deaza-dATP, dITP and hydroxymethyl-dUTP. Suitable Tm value-lowering substances are formanide, dimethyl sulphoxide and trimethylglycine. The methods improve the amplification of DNA. Also, DNA fragments which originated as RNA can be amplified without
                                  Human; Mcl-1 gene regulatory element; Mcl-1s/deltaTM variant; neuronal cell; tumour cell; apoptosis; therapy; cancer; psori diabetic retinopathy; corneal graft neovascularisation; neovascular glaucoma; epithelial condition; autoimmune diseas rheumatoid arthritis; systemic lupus erythematosus;
                                                                                                                                                                                                                                                                                                                                                                                               1014 CTTCCATGTAGAGGACCTAGA 1034
                                                                                                                                               Human Mcl-1 cDNA amplifying
                                                                                                                                                                                        10-AUG-2001
                                                                                                                                                                                                                                                                AAD06439 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    amplification
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Simple
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          21-OCT-1997;
14-AUG-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DNA amplification;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 1999-181059/15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (TAKI ) TAKARA SHUZO CO LTD.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WO9909213-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PCR primer P13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 10-AUG-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 27-MAY-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAX26539
                                                                                                                                                                                                                                                                                                                                                              21
                                                                                                                                                                                                                                                                                                                                                            CTTCCATGTAGAGGACCTAGA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            and accurate method for DNA amplification - uses cation in the presence of nucleotide analogues together with bund which lowers the Tm of double-stranded nucleic acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Kato I,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                the RNAs in sample.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Page 31;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           BP; 5 A; 5 C; 5 G; 6 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          97JP-0305016
97JP-0231885
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 98WO-JP03566
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mukai H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 nucleotide analogue; PCR primer; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         36pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                      0.5%;
100.0%;
                                                                                                                                                                                                                                                                  21
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            PCR primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BP.
                                                                                                                                                                                                                                                                                                                                                                                                                             0;
                                                                                                                                               RT-PCR primer, Mcl-1-F5.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Yamamoto
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 21;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 20;
. 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                 0;
                                                                                   cancer; psoriasis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                 Indels
                                                     disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                 0;
AC XXX AC
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RESULT 6
AAD06441/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local :
                                                                                                                                Homo
                                                                                                                                                                                            neuronal cell; tumour cell; apoptosis; therapy; cancer; psorias diabetic retinopathy; corneal graft neovascularisation; neovascular glaucoma; epithelial condition; autoimmune disease; rheumatoid arthritis; systemic lupus erythematosus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       expression in a cell e.g., neuronal cell or tumour cell, such that apoptosis of the cell is induced or cell viability is increased. The MCl-1 and its regulatory elements are used for treating pathological conditions which include cancer, diabetic retinopathy, corneal graft neovascularisation and neovascular glaucoma, epithelial conditions such as psoriasis, autoimmune diseases like rheumatoid arthritis, systemic lupus erythematosus, and neurodegenerative diseases. The present sequence is a RT-PCR primer Mcl-1-F5, used for the amplification of the human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present invention relates to Mcl-1 gene regulatory elements and t variant Mcl-1s/deltaTM. The anti-apoptotic Mcl-1 protein is encoded by exons 1, 2 and 3. The pro-apoptotic Mcl-1s/DeltaTM variant encoded by exons 1 and 3 is obtained due to alternative mRNA splicing. The Mcl-1 gene regulatory element is useful for modulating the Mcl-1 gene
14-JAN-2000; 2000WO-US00969.
                                                                                                                                                                                                                                                                Human; Mcl-1 gene regulatory element; Mcl-1s/deltaTM variant;
neuronal cell; tumour cell; apoptosis; therapy; cancer; psori
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAD06441 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Novel Mcl-1 gene regulatory of Mcl-1 polypeptide or its neuronal or tumor cells -
                                           25-MAY-2001.
                                                                                                                                                                         neurodegenerative
                                                                                                                                                                                                                                                                                                                                 Human
                                                                                                                                                                                                                                                                                                                                                                             10-AUG-2001
                                                                                                                                                                                                                                                                                                                                                                                                                           AAD06441;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2001-343812/36.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      16-NOV-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               14-JAN-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WO200136594-A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (DART-) DARTMOUTH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2; Page 75; 125pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Bingle CD,
                                                                                                                                                                                                                                                                                                                               cDNA amplifying RT-PCR primer,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            BP; 3 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2000WO-US00969
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                                                                                                                                                                           tis; systemic lupus erythematosus;
disease; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0.5%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             elements, useful for modulating expression variant which regulate apoptosis in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 21;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ×
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17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              22;
                                                                                                                                                                                                                                                                                                                                 Mc1-1-R7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0,:
                                                                                                                                                                                                                                                             cancer; psoriasis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 21;
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16-NOV-1999;

99US-0166113

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AAT03166/c
ID AAT031
XX AAT031
XX AAT031
XX O5-JUN
XX WC1-1;
KW WC1-1;
KW WC2-1;
KW Getect
XX Synthe
YN WO9528
YN WO9528
YN I2-APP
YN 12-APP
YN I3-APF
XX I3-APF
YN WPI;
PR 13-APF
YN WPI;
PR 13-APF
YN WPI;
PR 13-APF
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YN WPI;
PN WPI;
PN WPI;
PN WPI;
PN Detect
PN Detect
PN With F
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Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present invention relates to Mcl-1 gene regulatory elements and the variant Mcl-1s/deltaTM. The anti-apoptotic Mcl-1 protein is encoded by exons 1 2 and 3. The pro-apoptotic Mcl-1s/DeltaTM variant encoded by exons 1 and 3 is obtained due to alternative mRNA splicing. The Mcl-1 gene regulatory element is useful for modulating the Mcl-1 gene expression in a cell e.g., neuronal cell or tumour cell, such that apoptosis of the cell is induced or cell viability is increased. The Mcl-1 and its regulatory elements are used for treating pathological conditions which include cancer, diabetic retinopathy, corneal graft neovascularisation and neovascular glaucoma, peithelial conditions such as psoriasis, autoimmune diseases like rheumatoid arthritis, systemic lupus erythematosus, and neurodegenerative diseases. The present sequence is a RT-PCR primer Mcl-1-R7, used for the amplification of the human
                              Detection of novel proteins involved with proteins involved in apoptosis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1230 GAAAAGCAAGTGGCAAGAGGA 1250
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Novel Mcl-1 gene regulatory elements, useful for modulating expression of Mcl-1 polypeptide or its variant which regulate apoptosis in neuronal or tumor cells -
Example 1; Page 23; 62pp; English.
                                                                         WPI; 1995-373811/48
                                                                                                                                                                                                                                                      W09528497-A1.
                                                                                                                                                                                                                                                                                                             Mcl-1; Bax; apoptosis; cell death; regulation; Bcl-2; novel;
detection; ss.
                                                                                                                                                                                                                                                                                                                                                                                 05-JUN-1996 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Craig RW,
                                                                                                       Reed JC,
                                                                                                                                  (LJOL-) LA JOLLA CANCER RES FOUND
                                                                                                                                                                13-APR-1994;
                                                                                                                                                                                                                         26-OCT-1995
                                                                                                                                                                                                                                                                                  Synthetic
                                                                                                                                                                                                                                                                                                                                                     Human Mcl-1 gene reverse PCR primer.
                                                                                                                                                                                                                                                                                                                                                                                                                 AAT03166;
                                                                                                                                                                                                                                                                                                                                                                                                                                            AAT03166 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence
                                                                                                                                                                                           12-APR-1995;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GAAAAGCAAGTGGCAAGAGGA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CDNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity
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                                                                                                       Sato
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Page 75;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BP; 1 A;
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                                                                                                                                                              94US-0226876
                                                                                                                                                                                           95WO-US04600
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Pred. No.
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17;
                                            apoptosis
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                                            interaction
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                               The present invention describes antisense oligonucleotides which modulate the expression of novel anti-apoptotic bcl-2-related proteins. The antisense oligonucleotides can be used as therapeutic agents to prevent or delay inflammation or tumour formation by promoting apoptosis in human cells or tissues. They can also be used as research agents to establish the function of particular genes and as diagnostic agents in sandwich assays for detecting the level of novel anti-apoptotic bcl-2-related proteins in a sample. The antisense oligonucleotides given in the present invention were designed to target human Al and mcl-1 anti-apoptotic bcl-2-related protein nucleotide sequences. The present sequence represents an antisense oligonucleotide for the human mcl-1 nucleotide sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAT03164-T031366 are primers used for the amplification of the human Mcl-1 gene. The Mcl-1 protein is a Bcl-2 related protein, the Mcl-1 protein was expressed and used to investigate the interactions of Mcl-1 with other proteins known to be involved in apoptosis (excluding the Bax protein). Proteins detected using this method can act as upstream activators or downstream effectors of a cellular protein such as Bax which induces apoptosis. If the protein is a Bcl-2 related protein apoptosis levels are decreased due to the protein binding to and inactivating Bax.
                                                                                                                                                                                                                                                                                         Antisense oligonucleotides which modulate the expression of novel anti-apoptotic bcl-2-related proteins are useful for inducing apo and treating associated diseases e.g. cancer -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human; A1; anti-apoptotic; bcl-2-related protein; antisense inhibition;
mcl-1; apoptosis; cancer; antiinflammatory; cytostatic; tumour;
inflammation; diagnosis; phosphorothioate; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1182 GGACTCCAAGCTGTAACTTC 1201
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                                                                                                                                                                                                                                                           Claim 31;
                                                                                                                                                                                                                                                                                                                                                                                                Ackermann EJ, Marcusson
                                                                                                                                                                                                                                                                                                                                                                                                                               (ISIS-) ISIS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  07-JAN-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human mcl-1 anti-apoptotic bcl-2-related protein antisense oligo #20407
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Pred. No.
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51;
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Sequence 20 BP; 4 A; 6 C; 3 G; 7 T; 0 other;

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RESULT

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AAZ39084/c
ID AAZ39084 standard;
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Best Local
                                                                                                                                                                                                                                                       The present invention describes antisense oligonucleotides which modulate the expression of novel anti-apoptotic bcl-2-related proteins. The antisense oligonucleotides can be used as therapeutic agents to prevent or delay inflammation or tumour formation by promoting apoptosis in human cells or tissues. They can also be used as research agents to establish the function of particular genes and as diagnostic agents in sandwich assays for detecting the level of novel anti-apoptotic bcl-2-related proteins in a sample. The antisense oligonucleotides given in the present invention were designed to target human Al and mcl-1 anti-apoptotic bcl-2-related protein nucleotide sequences. The
                                                                                                                                                                                                Sequence 20 BP; 0 A;
                                                                                                                                                                                                                             present sequence represents an antisense oligonucleotide
human mcl-1 nucleotide sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 31; Column 33; 28pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Antisense oligonucleotides which modulate the expression of novel anti-apoptotic bcl-2-related proteins are useful for inducing apoptosis and treating associated diseases e.g. cancer -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             07-JAN-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human mcl-1 anti-apoptotic bcl-2-related protein antisense oligo #20408
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                                                                                                 636 CGCCAAGGACACAAAGCCAA 655
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                                                                   CGCCAAGGACACAAAGCCAA
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                                                                                                                                                                                                4 C; 7
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                                                                                                                                               Score 20;
Pred. No.
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Pred. No.
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51;
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AAZ39086/c
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В
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Best Local S
Matches 20
     Human; Al; anti-apoptotic; bcl-2-related protein; antisense inhibition; mcl-1; apoptosis; cancer; antiinflammatory; cytostatic; tumour; inflammation; diagnosis; phosphorothloate; ss.
                                                                            Human mcl-1
                                                                                                                                                                                                                                                                       2366 GGTGGATGGAGAGACATTTG 2385
                                                                                                                                                                                                                                                                                                                                                                                                              The present invention describes antisense oligonucleotides which modulate the expression of novel anti-apoptotic bcl-2-related proteins. The antisense oligonucleotides can be used as therapeutic agents to prevent or delay inflammation or tumour formation by promoting apoptosis in human cells or tissues. They can also be used as research agents to establish the function of particular genes and as diagnostic agents in sandwich assays for detecting the level of novel anti-apoptotic bcl-2-related proteins in a sample. The antisense oligonucleotides given in the present invention were designed to target human Al and mcl-1 anti-apoptotic bcl-2-related protein in uncleotide sequences. The present sequence represents an antisense oligonucleotide for the human mcl-1 nucleotide sequence.
                                                                                                                                           AAZ39086;
                                                                                                                                                                         AAZ39086 standard; DNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim
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mcl-1; apoptosis; cancer; antiinflammatory; cytostatic; tumour;
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ilarity 100.0%;
Conservative
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Pred. No.
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51;
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RESULT 10 AAZ39085/c

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Best Loc Matches

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RESULT 12
AAZ3907/C
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XX 29-FEH
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       or delay inflammation or tumour formation by promoting apoptosis in human cells or tissues. They can also be used as research agents to establish the function of particular genes and as diagnostic agents in sandwich assays for detecting the level of novel anti-apoptotic bcl-2-related proteins in a sample. The antisense oligonucleotides given in the present invention were designed to target human Al and mcl-1 anti-apoptotic bcl-2-related protein nucleotide sequences. The present sequence represents an antisense oligonucleotide for the human mcl-1 nucleotide sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Antisense oligonucleotides which modulate the expression of novel anti-apoptotic bcl-2-related proteins are useful for inducing apoptosis and treating associated diseases e.g. cancer - \frac{1}{2}
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            07-JAN-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present invention describes antisense oligonucleotides which modulate the expression of novel anti-apoptotic bcl-2-related proteins. The antisense oligonucleotides can be used as therapeutic agents to prevent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2000-061908/05
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 31; Column 33; 28pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Ackermann
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (ISIS-) ISIS
     (ISIS-) ISIS PHARM INC.
                                                      07-JAN-1999;
                                                                                                      07-JAN-1999;
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                                                                                                                                                                                                                  US6001992-A.
                                                                                                                                                                                                                                                                                         Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                Human; A1; anti-apoptotic; bcl-2-related protein; antisense inhibition;
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human mcl-1 anti-apoptotic bcl-2-related protein antisense oligo #20421
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         29-FEB-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAZ39087;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAZ39087 standard;
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                                                                                                                                                                                                                                                                                                                                                  inflammation;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   l Similarity 100.0%;
20; Conservative
                                                                                                                                                                                                                                                                                                                                                                            apoptosis; cancer; antiinflammatory;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            BP; 4 A; 7 C; 3 G; 6 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Marcusson
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0.5%;
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                                                                                                                                                                                                                                                                                                                                                  phosphorothioate;
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DB 21;
                                                                                                                                                                                                                                                                                                                                                        SS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Dean
                                                                                                                                                                                                                                                                                                                                                                         cytostatic; tumour;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The present invention describes antisense oligonucleotides which modulate the expression of novel anti-apoptotic bcl-2-related proteins. The antisense oligonucleotides can be used as therapeutic agents to prevent or delay inflammation or tumour formation by promoting apoptosis in human cells or tissues. They can also be used as research agents to establish the function of particular genes and as diagnostic agents in sandwich assays for detecting the level of novel anti-apoptotic bcl-2-related proteins in a sample. The antisense oligonucleotides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Antisense oligonucleotides which modulate the expression of novel anti-apoptotic bcl-2-related proteins are useful for inducing apoptosis and treating associated diseases e.g. cancer \cdot
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       given in the present invention were designed to target human Al and mcl-1 anti-apoptotic bcl-2-related protein nucleotide sequences. The present sequence represents an antisense oligonucleotide for the human mcl-1 nucleotide sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                       Human mcl-1 anti-apoptotic bcl-2-related protein antisense oligo #20404.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         3391 TAAAAGCCTGTCTGCCAAAT 3410
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 31; Column 33;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2000-061908/05
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                                                                                                                                                                                                                                                                                                                                    Homo
                                                                                                                                                                                                                                                                                                                                                                                               Human; Al; anti-apoptotic; bcl-2-related protein; mcl-1; apoptosis; cancer; antiinflammatorv: cvtost
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAZ39090;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAZ39090 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 20 BP; 5 A; 3 C; 5 G; 7 T; 0 other;
                                                        Antisense oligonucleotides which modulate the expression of novel anti-apoptotic bcl-2-related proteins are useful for inducing apoptosis and treating associated diseases e.g. cancer - \,
                                                                                                                         WPI;
                                                                                                                                                       Ackermann
                                                                                                                                                                                  (ISIS-) ISIS
                                                                                                                                                                                                                 07-JAN-1999;
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                                                                                                                                                                                                                                                                                                                                                   Synthetic
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                                                                                                                                                                                                                                                                                                                                                                                inflammation;
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                                                                                                                                                                                                                                                                                                                                   sapiens
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                                                                                                                         2000-061908/05
                                                                                                                                                                                                                                                                                                                                                                               apoptosis; cancer; antiinflammatory; cytostatic;
mation; diagnosis; phosphorothioate; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity
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                                                                                                                                                       EJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                    PHARM INC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0.5%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 20; DB 21;
Pred. No. 51;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Bennett
                                                                                                                                                        Bennett CF,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CF,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Dean
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                                                                                                                                                                                                                                                                                                                                                                                                               antisense
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                                                                                                                                                                                                                                                                                                                                                                                                                 inhibition
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The present invention describes antisense oligonucleotides which modulate

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Column 33;

28pp; English.

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RESULT
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Best L
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                                                                                      The present invention describes antisense oligonucleotides which modulate the expression of novel anti-apoptotic bcl-2-related proteins. The antisense oligonucleotides can be used as therapeutic agents to prevent or delay inflammation or tumour formation by promoting apoptosis in human cells or tissues. They can also be used as research agents to establish the function of particular genes and as diagnostic agents in sandwich assays for detecting the level of novel anti-apoptotic cl-2-related proteins in a sample. The antisense oligonucleotides are the market and a sample of the antisense oligonucleotides.
                                                           given
mcl-1
                                                                                                                                                                                                                                                                                                                                                                     Antisense oligonucleotides which modulate the expression of novel anti-apoptotic bcl-2-related proteins are useful for inducing apopand treating associated diseases e.g. cancer -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           the expression of novel anti-apoptotic bcl-2-related proteins. The antisense oligonucleotides can be used as therapeutic agents to prevent or delay inflammation or tunour formation by promoting apoptosis in human cells or tissues. They can also be used as research agents to establish the function of particular genes and as diagnostic agents in sandwich assays for detecting the level of novel anti-apoptotic bcl-2-related proteins in a sample. The antisense oligonucleotides given in the present invention were designed to target human Al and mcl-1 anti-apoptotic bcl-2-related protein nucleotide sequences. The present sequence represents an antisense oligonucleotide for the branch man antial management.
                                                                                                                                                                                                                                                                                                                                  Example 12; Column 33;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (ISIS-) ISIS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      07-JAN-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; Al; anti-apoptotic; bcl-2-related protein; antisense inhibition;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  inflammation;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2000-061908/05
in the present invention were designed to target human Al and anti-apoptotic bcl-2-related protein nucleotide sequences. The nt sequence represents an antisense oligonucleotide for the mcl-1 nucleotide sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      mcl-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              l Similarity
20; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          apoptosis;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         anti-apoptotic bcl-2-related protein antisense oligo #20405
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BP; 5 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           nucleotide sequence
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ilarity 100.0%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             diagnosis; phosphorothioate; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Marcusson
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        cancer;
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                                                                                                                                                                                                                                                                                                                          28pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             C; 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  EG,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     antiinflammatory;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                G; 3 T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 20;
Pred. No.
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Search completed:

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RESULT 15
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Matches 20
                                                                                                                                                                                                                                                                 antisense oligonucleotides can be used as therapeutic agents to prevent or delay inflammation or tumour formation by promoting apoptosis in human cells or tissues. They can also be used as research agents to establish the function of particular genes and as diagnostic agents in sandwich assays for detecting the level of novel anti-apoptotic bcl-2-related proteins in a sample. The antisense oligonucleotides given in the present invention were designed to target human Al and mcl-1 anti-apoptotic bcl-2-related protein nucleotide sequences. The present sequence represents an antisense oligonucleotide for the human mall and present sequence represents an antisense oligonucleotide for the
                                                                                                                                                                                                      Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present invention describes antisense oligonucleotides which modulate the expression of novel anti-apoptotic bel-2-related proteins. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Antisense oligonucleotides which modulate the expression of novel anti-apoptotic bcl-2-related proteins are useful for inducing apo and treating associated diseases e.g. cancer -
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Job time : 579.329 secs

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AU240558 AU240558 AU009945 AU009945 AZ482026 1M0306K18 BH849581 SALK_0699 AL765626 Arabidops BH011458 BG02105-5

AZ787019 2M0032B19 AZ323085 1M0044J02 AL761633 Arabidops AZ809484 2M0073F04

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Minimum
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AI370619
AI370619.1
               Tumor Gene Index
Unpublished (1997)
Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further information.
Trace considered overall poor quality
Insert Length: 431 Std Error: 0.00
Seq primer: -40UP from Gibco
High quality sequence stop: 1.
                                                                                                                                                                                                A370619 43 bp mRNA linear EST 16-FEB-1
ta40f05.x1 Soares_total_fetus_Nb2HFB_9w Homo sapiens cDNA clone
IMAGE:2046561 3' similar to SW:BFLL HUMAN Q16548 BCL2-RELATED
PROTEIN A1 ; contains element TAR1 repetitive element ;, mRNA
                                                                                                    Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 43) NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap. NCI-CGAP institute, Cancer Genome Anatomy Project (CGAP), Mational Cancer Institute, Cancer Genome Anatomy Project (CGAP),
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AW332749 S12E1 AGS AZ346753 1M0082E11 BH863143 SALK 0932 AU254252 AU254252 AU254252 AU254252 AZ331642 1M0059P11 AZ469379 1M0282P14 AZ663912 1M0543J18

EST 16-FEB-1999

D45819 HUMGS03038 AU257038 AU257038 AU259737 AU259737

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Zea mays.
Zea mays
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Possible ligation site of ends
                                                                                                                                                                                                                                                                                                                                                                                                             855 California Ave, Palo Alto, CA 94304, USA
Tel: 650 723 2227
Fax: 650 725 8221
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Stanford University
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Unpublished (2001
Contact: Walbot V
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Maize genomic sequences found using engineered RescueMu transposon
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                                                                                                                                                                                                                                                                                                      transposon-tagged.
Location/Qualifiers
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/note="Organ: leaf; Vector: RescueMu (engineered from pBlueScript backbone); Site_1: BanHI; Site_2: BglII; BBHEIS at 4.9 kb, modified maize Mu transposon designed to allow plasmid rescue from total genomic DNA. Mu elements insert preferentially into transcription units. For more information on RescueMu go to the web site www.zmdb.iastate.edu and follow the links for RescueMu.' Grid I was grown at Berkeley in 2001. DNA wa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /cultivar="mixed background W23/A188/B73"
/db_xref="taxon:4577"
/clone_lib="1008 - RescueMu Grid I"
/tissue_type="leaf"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /note="Vector: pT7T3D-Pac (Pharmacia) with a modified
polylinker; Site_1: Not I; Site_2: Eco RI; 1st strand cDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /clone="IMAGE:2046561"
/clone_lib="Soares_total_fetus_Nb2HF8_9w"
/dev_stage="8-9 weeks"
/lab_host="bH10B"
                                                                                                                                                        /lab_host="DH10B"
                                                                                                                                                                           /dev_stage="adult"
                                                                                                                                                                                                                                                                   /organism="Zea mays"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /organism="Homo sapiens"
/db_xref="taxon:9606"
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Pred. No.
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. 0.00029;
thes 0;
                                                                                                                                                                                                                                                                                                                                                       by 2 different endonucleases. sequence from source sequence.
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AUTHORS
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REFERENCE AUTHORS

COMMENT

FEATURES

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VERSION KEYWORDS

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Matches Query Match

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JOURNAL
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                                                                                                                   Query Match
                                                                                                 Best Local Similarity
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Best Local :
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42
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TGGGAAAATGGCTTT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1-3 yamada-oka, Suita, Osaka Pref. 565, Japan
Tel: 06-877-5111(ex.3315)
Email: kousaku@imcb.osaka-u.ac.jp
Email: kousaku@imcb.osaka-u.ac.jp
Email: kousaku@imcb.osaka-u.ac.jp
Email: kousaku@imcb.osaka-u.ac.jp
For the abundance information of clones with this sequence in this
For the abundance information of clones with this sequence in this
Ibrary and as well as in other 3'-directed libraries, see '
http://www.imcb.osaka-u.ac.jp/bodymap'. The sequences of the clones
represented by this GS sequences is also found there.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     l Similarity
16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Osaka University
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Contact: Okubo, K
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Okubo,K.

BodyMap; human gene expression database Unpublished (1995)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
1 (bases 1 to 43)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens
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                                                                              Conservative
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                                                                                                                                                                                                                                      /note="Organ: blood; Vector: 1-gt-11; Site_1: Eco-RI; Monocytes were prepared from blood by ficoll-hypaque, percoll and T cell rosetting purification steps (purity: 96 %). mRNA was prepared from activated monocytes from a patient with rheumatoid arthritis. mRNA was reverse transcribed with MuLY. Using Eco-RI linkers cDNA was cloned into 1-gt-11 vector arms. The cDNA library was screened by differential hybddization using radioactively
                                                                                                                                                                                monocytes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      extracted from leaf punches, double digested using BamHI and BgLII, and ligated to form circular plasmids. DHIOB cells were transformed and then screened on LB plates with ampicillin.
                                                                                                                                                                                                                       marked ss-cDNA from activated and non-activated
                                                                                                                                                                                                                                                                                                                                                                                                     /db_xref="taxon:9606"
/clone_lib="Human adult (K.Okubo)"
/dev_stage="adult"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /organism="Homo sapiens"
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28
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                                                                              0;
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                                                                                               Score 15; DB 14;
Pred. No. 4.7e+03;
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. 1.3e+03;
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REFERENCE
AUTHORS
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                                                                                            719 ACCTTCCAGAGTTA 732
                                                                                                                                                       Local Similarity
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                                                     ACCTTCCAGAGTTA 20
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AZ439313. GI:10563326
GSS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Unpublished (2000)
Contact: Robert B. Weiss
University of Utah
Rm. 308, Blomedical Polymers Research Bldg.,
Balla, USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D.,Weiss,R. and wright,D.,Weiss,R.
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Insert Length: 10000 Std Error: 0.00
Plate: 0229 row: J column: 20
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1 (bases 1 to 26)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      eq primer: CACACAGGAAACAGCTATGACC lass: plasmid ends igh quality sequence stop: 26.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             801 585 5606
801 585 7177
                                                                                                                                                                                                                            (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XLIO-Gold (Stratagene) cells and selected for ampicillin resistance."
                                                                                                                                                                                                                                       12
                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-" /note="Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /organism="Mus musculus"
/strain="C57BL/6J"
/db_xref="taxon:10090"
/clone="UUGC1M0229J20"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /clone_lib="Mouse 10kb plasmid UUGC1M library"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Laboratory Mouse DNA Resource
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                                                                                                                                                       1.8%;
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                                                                                                                                                     Score 14; DB 17;
Pred. No. 1.4e+04
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                                                                                                                                     Mismatches
                                                                                                                                                       1.4e+04;
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Salk Institute Genomic Analysis Laboratory (SIGNAL)
The Salk Institute for Biological Studies
10010 N. Torrey Pines Road, La Jolla, CA 92037, USA
Tel: 858 453 4100 x1752
Fax: 858 558 6379
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Trypanosoma brucei
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Alonso, J.M., Leisse, T.J., Barajas, P., Chen, H., Cheuk, R., Gadrinab, C., Jeske, A., Karnes, M., Kim, C.J., Parker, H., Prednis, L., Shinn, I, Zimmerman, J. and Ecker, J.R.
A Sequence-Indexed Library of Insertion Mutations in the
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                                                                          Hall,N., Bowman,S., Lennard,N.J., Doggett,J., Atkin,R., Chillingworth,C., Ormond,D., Harris,B., El-Sayed,N., Hou,L., Melville,S.E., Rajandream,M.A. and Barrell,B.G.
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Direct Submission Submitted (10-DEC-2000) Trypanosoma brucei genome sequencing
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/clone=lib="Arabidopsis thaliana TDNA insertion lines"
/clone_lib="Arabidopsis thaliana TDNA insertion lines
/note="pck was performed on Arabidopsis thaliana lines
each of which contains one or more TDNA insertion
elements. The resultant fragment for each line was
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                                                                                                                                                    Email: ddunn@genetics.utah.edu
Insert Length: 1000 Std Error:
Plate: 0032 row: B column: 19
Seg primer: CACAAGGAAACAGCTATGACC
Class: plasmid ends
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              26 bp DNA linear GSS 16-FEB-200032B19R Mouse 10kb plasmid UUGC1M library Mus musculus genomic clone UUGC2M0032B19 R, DNA sequence.
AZ787019
                                                                                                               High quality sequence stop: 26.
Location/Qualifiers
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Rockville, MD. Genomic DNA isolated from a cloned population of
Trypanosoma brucei (TREUD27/4 GUTat 10.1) was mechanically sheared
to give a tight size distribution (
4 kb). The v + i method used for the library construction is
described in detail in Smith, H. and Venter, J.C. (Making small
insert libraries for whole genome shotgun sequencing projects. In
Genome Sequencing: A Practical Approach, eds. M. Vaudin and B.
Barrell, Oxford University Press, 1999).
Detail: nelsayed@tigr.org
                                                                                                                                                                                                                                                 Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                        University of Utah Genome Center University of Utah
                                                                                                                                                                                                                                                                                                                                                                               plasmid inserts
Unpublished (2000)
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                                                                                                                                                                                                                                                                                                                                                                Contact: Robert B.
                                                                                                                                                                                                                                                                                                                                                                                                               Mouse whole genome scaffolding with paired end reads from 10kb
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           house mouse
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/db_xref="taxon:10090"
/clone="UUGC2M0032B19"
/clone_lib="Mouse 10kb plasmid UUGC1M library"
                                                          /organism="Mus musculus"
/strain="C57BL/6J"
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/strain="TREU927"
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                                                                                                                                                                                                                                             Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                          Contact: Robert B.
                                                                                                                                                                                                                                                                                                                                                                         Unpublished (2000)
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1 (bases 1 to 29)
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Plate: 0044 row: J column: 02
Seq primer: CGTTGTAAAACGACGGCCAGT
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University of Utah
Rm. 308, Biomedical Polymers Research Bldg.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1M0044J02F Mouse 10kb plasmid UUGC1M clone UUGC1M0044J02 F, DNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AZ323085.1 GI:10377460
                                                                                                                                                                      quality sequence stop: 29.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                       ddunn@genetics.utah.edu
Length: 10000 Std Error:
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/clone_lib="Mouse 10kb plasmid UUGC1M library"
                                 /clone="UUGC1M0044J02"
                                                         /db_xref="taxon:10090"
                                                                                 /organism="Mus musculus"
/strain="C57BL/6J"
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/lab_host="E. Coli strain XL10-Gold, Tl-resistant, F-"
/lab_host="E. Coli strain XL10-Gold, Tl-resistant, F-"
/note="Vector: PWD42nv; Purified genomic DNA from M.
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Pred. No.
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GSS
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AL761633
                                                                              Direct Submission
Submitted (17-UN-2002) Weisshaar B., Max-Planck-Institut fuer Submitted (17-UN-2002) Weisshaar B., Max-Planck-Institut fuer Zuechtungsforschung, Carl-von-Linne-Weg 10, Koeln, 50829, Germany This sequence is recovered from the left border of the T-DNA. It indicates an insertion within the locus defined by clone MPF21. The sequences are generated at the MPI for Plant Breeding Research in the context of the GABI-Kat project. GABI-Kat is part of the German Plant Genomics program designated 'GABI'. Information on line availability can be found at:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A pipeline for automated high-throughput generation of FSTs (flanking sequence tags) from Arabidopsis thaliana T-DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Strizhov, N., Li, Y., Rosso, M., Viehoever, P., Dekker, K., Saedler, H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; Rosidae; eurosids II; Brassicales; Brassicaceae; Arabidopsis.
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                                                                                                                                                                                                                                                                                                                                                                                                                                Unpublished
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Unpublished
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/organism="Arabidopsis thaliana"
                                                           Location/Qualifiers
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
Rm. 308, Biomedical Polymers Rese
84112, USA
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13; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D.,Weiss,R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AZ809484 31 bp DNA linear GSS 20-FEB-200 2M0073F04R Mouse 10kb plasmid UUGC1M library Mus musculus genomic clone UUGC2M0073F04 R, DNA sequence.
                                                                                                                                                                                                                                                                                                                                                                      Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: (
Plate: 0073 row: F column: 04
Seg primer: CACACAGGAAACAGCTATGACC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Unpublished (2000)
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/db_xref="taxon:3702"
/db_xref="taxon:3702"
/clone="GK-227B05-014266"
/clone=lib="Arabidopsis thaliana T-DNA insertion lines"
/note="PCR was performed on DNA from Arabidopsis thaliana
plants (T1) which were transformed with the T-DNA from
vector pAC161. The lines contain one or more T-DNA
insertions. The DNA fragment(s) resulting from the PCR
were directly sequenced to determine the genomic sequence
flanking the insertion. Sequences displaying significant
similarity to the A. thalaina nuclear genome sequence were
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     processed for submission. T-DNA derived sequences were removed" 7 g 8 t
Laboratory Mouse DNA Resource (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA
                                                                                       /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-" /note="Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson
                                                                                                                                                         /clone_lib="Mouse 10kb plasmid UUGC1M library"
/sex="Male"
                                                                                                                                                                                               /organism="Mus musculus"
/strain="C57BL/6J"
/db_xref="taxon:10090"
/clone="UUGC2M0073F04"
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Pred. No.
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les 13; Conserv
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                                TCAACAGCTTCAA 38
TCAACAGCTTCAA 28
                                                                                                                                                                                                                                                                                                                                                                                                                                    Department of Biological Sciences
Graduate School of Science, University of Tokyo
Hongo 7-3-1, Bunkyo-ku, Tokyo 113-0033, Japan
                                                                                                                                                                                                                                                                                                                                        Email: naruse@biol.s.u-tokyo.ac.jp
This clone was isolated from UV ir:
library (OLC) 5' end sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Unpublished (2001)
Contact: Kiyoshi Naruse
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Naruse,K., Mitani,H. and Tanaka,M. Medaka EST Project in University of Tokyo (2001)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Actinopterygii; Neopterygii; Teleostei; Euteleostei; Neoteleostei; Acanthomorpha; Acanthopterygii; Percomorpha; Atherinomorpha; Beloniformes; Adrianichthyldae; Oryziinae; Oryzias.

1 (bases 1 to 33)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Oryzias latipes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AU240558 UV irradiated OLHNI cell line cDNA library latipes cDNA clone OLc21.11d, mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                        Tel: 81-3-5841-4443
                                                                                                                                                                                                                                                                                                                                                                                                        Fax: 81-3-5841-4410
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Japanese medaka.
                                                                           Conservative
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                                                                                                                                                                                        (OLc)"
                                                                                                                                                                                                     /db_xref="taxon:8090"
/clone="OLc21.11d"
/clone=lib="DV irradiated OLHNI cell line cDNA library
                                                                                                                                                                                                                                                                /organism="Oryzias latipes'
/strain="HNI"
                                                                                                                                                                                                                                                                                                                          Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         8 ATATATTTACAGG 20
                plasmid inserts
Unpublished (2000)
Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
Rm. 308, Biomedical Polymers Research Bldg.,
                                                                                                                                                                                                            Mus musculus

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

Bukaryota; Metazoa; Chordata; Sciurognathi; Muridae; Murinae; Mus

Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus

1 (bases 1 to 36)

Rarher.M., Beacorn,T., Duval,B., Hamil,C.,,
                                                                                                                                                                                                                                                                                                                                 AZ482026.1
GSS.
                                                                                                                                                                                            Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Rei
                                                                                                                                                                                                                                                                                                                                                                                    AZ482026 36 bp DM
1M0306K18R Mouse 10kb plasmid UUGC1M
clone UUGC1M0306K18 R, DNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Genome Research Group
National Institute of Radiological Sciences
9-1, Anagawa-4-chome, Inage-ku, Chiba, Chiba
                                                                                                                                       Mouse whole genome scaffolding with paired
                                                                                                                                                                           Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly
,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A.
                                                                                                                                                                                                                                                                                                                                                                          AZ482026
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AU009945 Schizosaccharomyces pombe late log phase cDNA
Schizosaccharomyces pombe cDNA clone spc00705, mRNA sequence.
                                                                                                                                                             and Wright,D.,Weiss,R.
                                                                                                                                                                                                                                                                                                                   house mouse.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Morimyo, M. and Mita, K. Identification of expressed sequence
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1 (bases 1 to 35)
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308,
308,
USA
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Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               D
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /note="Vector: M13mp19; The cDNA library of Schizosaccharomyces pombe was prepared by cloning cDNA into the Smal site of M13mp19 DNA and the direction of sequences was not always from 5' to 3'. The cDNA data Schizosaccharomyces pombe are available for searching the World Wide Web. (URL, http://www.nirs.go.jp)" 8 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /db_xref="taxon:4896"
/clone="spc00705"
/clone_lib="Schizosaccharomyces pombe late log phase cDNA"
/sex="h_minus"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /strain="972"
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Pred. No. 5.7e+04;
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                                                                                                                                         from 10kb
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RESULT 14
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Matches 13
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Local Similarity 100.0%; local Similarity 00.0%; last the conservative 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TTTTGTTGCGGAG 450
                                                                                                                                                     Alonso,J.M., Leisse,T.J., Barajas,P., Chen,H., Cheuk,R., Gadrinab,C., Jeske,A., Karnes,M., Kim,C.J., Parker,H., Prednis,L., Shinn,P., Zimmerman,J. and Ecker,J.R.
A Sequence-Indexed Library of Insertion Mutations in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BH849581 36 bp DNA linear GSS 13-JUN-2002 SALK_069905.21.55.x Arabidopsis thaliana TDNA insertion lines Arabidopsis thaliana genomic clone SALK_069905.21.55.x, DNA
                       Arabidopsis Genome
Unpublished (2001)
Contact: Joseph R. Ecker
Salk Institute Genomic Analysis Laboratory (SIGnAL)
The Salk Institute for Biological Studies
                                                                                                                                                                                                                                                       Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnollophyta; eudicotyledons; core eudicots; Rosidae; eurosids II; Brassicales; Brassicaceae; Arabidopsis.

1 (bases 1 to 36)
                                                                                                                                                                                                                                                                                                                                                                             Arabidopsis thaliana
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Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (9144732114(gb)AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
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N. Torrey Pines Road,
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Length: 10000 Std Error: 0.00
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/clone="UUGC1M0306K18"
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/strain="C57BL/6J"
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                                               Zuechtungsforschung, Carl-von-Linne-Weg 10, Koeln, 50829, Germany This sequence is recovered from the left border of the T-DNA. It indicates an insertion close to or within gene At1929750. The sequences are generated at the MPI for Plant Breeding Research in the context of the GABI-Kat project. GABI-Kat is part of the Germa Plant Genomics program designated 'GABI'. Information on line availability can be found at:

http://www.mpiz-koeln.mpg.de/GABI-Kat/.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    thale cress.
Arabidopsis thaliana
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots;
Rosidae; eurosids II; Brassicales; Brassicaceae; Arabidopsis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 Similarity
13; Conserv
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Direct Submission
Submitted (17-JUN-2002) Weisshaar B., Max-Planck-Institut fuer
                                                                                                                                                                                                                                                                                                                                         Rosso,M., Strizhov,N., Li,Y., Reiss,B., Dekker,K. and Weisshaar,B. A new Arabidopsis thaliana T-DNA mutagenised population (GABI-Kat)
                                                                                                                                                                                                                                                                                                                                                                                                                                         A pipeline for automated high-throughput generation of FSTs (flanking sequence tags) from Arabidopsis thaliana T-DNA
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/organism="Arabidopsis thaliana"
/strain="Columbia 0"
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7 TCATATTTTGTTG 19
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/clone="GK-026F02-013759"
/clone="GK-026F02-013759"
/clone="lib="Narabidopsis thaliana T-DNA insertion lines"
/note="PCR was performed on DNA from Arabidopsis thaliana plants (T1) which were transformed with the T-DNA from vector pAc106. The lines contain one or more T-DNA insertions. The DNA fragment(s) resulting from the PCR were directly sequenced to determine the genomic sequence flanking the insertion. Sequences displaying significant similarity to the A. thalaina nuclear genome sequence were processed for submission. T-DNA derived sequences were removed"
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NAME: Imbra, Richard J.
REGISTRATION NUMBER: 37,643
REFERENCE/DOCKET NUMBER: FP-L.
TELECOMMUNICATION INFORMATION:
TELEPHONE: (619) 535-9001
TELEPHONE: (619) 535-9049
INFORMATION FOR SEQ ID NO: 1:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 3, Application US/08607269 Patent No. 5702897
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Best Local Similarity 100.0%;
Matches 21; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Reed, John C.
APPLICANT: Sato, Takaaki
TITLE OF INVENTION: Interaction of Proteins Involved in a
TITLE OF INVENTION: Cell Death Pathway
NUMBER OF SEQUENCES: 29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURENT APPLICATION UNMBER: PCT/US95/04600
FILING DATE: 12-APR-1995
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: LA JOLLA CANCER RESEARCH FOUNDATION
TITLE OF INVENTION: Interaction of Proteins Inv
TITLE OF INVENTION: a Cell Death Pathway
NUMBER OF SEQUENCES: 29
CORRESPONDENCE ADDRESS:
ADDRESSEE: Campbell and Flores
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
LENGTH: 21 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS:
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STREET
                                                                                                                                                                                                                           COUNTRY: UZIP: 92122
                                                                                                                                                                                                                                                                                                                                            ADDRESSEE: Campbell and Flores
STREET: 4370 La Jolla Village Drive, Suite
CITY: San Diego
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LENGTH: 21 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STATE: California
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                                                                                                                                                                                                                                                                                                        California
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                                                                                                                                                                                                                                                                    USA
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Pred. No.
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GENERAL INFORMATION: APPLICANT: Ackerman

APPLICANT: Dean, Nicholas M.

Bennett,

Frank

Ackermann, Elizabeth J.

APPLICANT:

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US-09-226-568-20/c
; Sequence 20, Application US/09226568
; Patent No. 6001992
                                                                                                                                                                                                                                                                   ; OTHER INFORMATION: Description of Artificial Sequence: antisense ; OTHER INFORMATION: sequence US-09-226-568-19
                                                            RESULT 5
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                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Ackermann, Elizabeth J.
APPLICANT: Bennett, C. Frank
APPLICANT: Bennett, C. Frank
APPLICANT: Dean, Nicholas M.
APPLICANT: Marcusson, Eric G.
TITLE OF INVENTION: Antisense Modulation of No. 6001992el Anti-apoptotic
TITLE OF INVENTION: Bcl-2-Related Proteins
FILE REFERENCE: ISPH-0337
FULE REFERENCE: ISPH-0337
CURRENT APPLICATION NUMBER: US/09/226,568
CURRENT FILING DATE: 1999-01-07
NUMBER OF SEQ ID NOS: 39
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 19, Application US/09226568 Patent No. 6001992
                                                                                                                                                                                          Matches
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                                                                                                                                                                                                            Best Local Similarity
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INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                          ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                  TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                  ENGTH:
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ATTORNEY/AGENT INFORMATION:
NAME: Campbell, Cathryn A.
REGISTRATION NUMBER: 31,81
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        REFERENCE/DOCKET NUMBER: P-LJ 9882
TELECOMMUNICATION INFORMATION:
TELEPHONE: (619) 535-9001
TELEPAX: (619) 535-8049
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION DATA:
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                                                                                                                   20 GGCGACTGGCAATGTTTGGC 1
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STRANDEDNESS: singl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TOPOLOGY:
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100.0%; Pr
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100.0%; Pred. No. 9.8;
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Pred. No.
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; OTHER INFORMATION:
US-09-226-568-21
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US-09-226-568-21/c
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; OTHER INFORMATION: sequence
US-09-226-568-20
                                                                                                                                                                                           RESULT 7
US-09-226-568-22/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION NUMBER: US/09/226,568
CURRENT FILING DATE: 1999-01-07
NUMBER OF SEQ ID NOS: 39
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 21
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CURRENT FILING DATE: 1999-01-07
NUMBER OF SEQ ID NOS: 39
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 20
LENGTH: 20
                                                                                                                                         GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local :
                                                                                                                                                          Sequence 22, Application US/09226568 Patent No. 6001992
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Best Local Similarity
             APPLICANT: Bennett, C. Frank
APPLICANT: Dean, Nicholas M.
APPLICANT: Marcusson, Eric G.
TITLE OF INVENTION: Antisense Modulation of No.
TITLE OF INVENTION: bcl-2-Related Proteins
FILE REFERENCE: ISPH-0337
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Ackermann, Elizabeth J.
APPLICANT: Bennett, C. Frank
APPLICANT: Dean, Cicholas M.
APPLICANT: Dean, Cicholas M.
APPLICANT: Marcusson, Eric G.
TITLE OF INVENTION: Antisense Modulation of No. 6001992el Anti-apoptotic
TITLE OF INVENTION: bcl-2-Related Proteins
FILE REFERENCE: ISPH-0337
CURRENT APPLICATION NUMBER: US/09/226,568
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TITLE OF INVENTION: Antisense Modulation of No. 6001992el Anti-apoptotic TITLE OF INVENTION: bc1-2-Related Proteins
FILE REFERENCE: ISPH-0337
                                                                                                                       APPLICANT: Ackermann,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: DNA ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FEATURE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH: 20
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Local Similarity 100.0%;
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                                                                                                                       Elizabeth J
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Pred. No.
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Pred. No
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9.8;
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                                                6001992el Anti-apoptotic
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; TYPE: DNA;
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Description of Artificial Sequence: antisense
; OTHER INFORMATION: sequence
US-09-226-568-22
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                                                                                                                                                                                                                                            GENERAL INFORMATION:
TITLE OF INVENTION: Antisense Modulation of No. 6001992el Anti-apoptotic TITLE OF INVENTION: bol-2-Related Proteins
FILE REFERENCE: ISPH-0337
CURRENT APPLICATION NUMBER: US/09/226,568
CURRENT FILING DATE: 1999-01-07
NUMBER OF SEQ ID NOS: 39
SOFTWARE: PATENTIN Ver. 2.0
SEQ ID NO 24
LENGTH: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 23
LEUGTH: 20
TYPE: DNA
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NUMBER OF SEQ ID NOS: 39
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 22
LENGTH: 20
                                                                                                                                                                                                                                                            Sequence 24, Application US/09226568 Patent No. 6001992
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
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Best Local
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                                                                                                                                                                   APPLICANT: Ackermann, Elizabeth J.
APPLICANT: Bennett, C. Frank
APPLICANT: Dean, Nicholas M.
APPLICANT: Marcusson, Eric G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION: Antisense Modulation of No. 6001992el Anti-apoptotic TITLE OF INVENTION: bcl-2-Related Proteins FILE REFERENCE: ISPH-0337
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APPLICANT: Dean, Nicholas M.
APPLICANT: Marcusson, Eric G.
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CURRENT FILING DATE: 1999-01-07
NUMBER OF SEQ ID NOS: 39
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                                                                                                                                                                                                                                                                                                                                                                                                                 636 CGCCAAGGACACAAAGCCAA 655
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APPLICANT: Dean, Nicholas M.
APPLICANT: Marcusson, Eric G.
TITLE OF INVENTION: Antisense Modulation of No. 60
TITLE OF INVENTION: Antisense Modulation of No. 60
TITLE OF INVENTION: bcl-2-Related Proteins
FILE REFERENCE: ISPH-0337
CURRENT APPLICATION NUMBER: US/09/226,568
CURRENT FILING DATE: 1999-01-07
NUMBER OF SEQ ID NOS: 39
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 26
LENGTH: 20
TYPE: DNA
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APPLICANT: Marcusson, Eric G.
APPLICANT: Marcusson, Eric G.
TITLE OF INVENTION: Antisense Modulation of No. 60:
TITLE OF INVENTION: Antisense Modulation of No. 60:
TITLE OF INVENTION: bcl-2-Related Proteins
FILE REFERENCE: ISPH-0337
CURRENT APPLICATION NUMBER: US/09/226,568
CURRENT FILING DATE: 199-01-07
NUMBER OF SEQ ID NOS: 39
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 25
LENGTH: 20
TYPE: DNA
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APPLICANT: Ackermann, Elizabeth J.
APPLICANT: Bennett, C. Frank
                                                                                                                                                                                                                                                                                                                                              Sequence 26, Application US/09226568
Patent No. 6001992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 0.5
Best Local Similarity 100.
Matches 20; Conservative
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OTHER INFORMATION: Description of Artificial Sequence: antisense OTHER INFORMATION: sequence
                                             ORGANISM: Artificial Sequence
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ORGANISM: Artificial Sequence
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Pred. No.
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APPLICANN: Ackermann, Elizabeth J.
APPLICANN: Bennett, C. Frank
APPLICANT: Bennett, C. Frank
APPLICANT: Bennett, C. Frank
APPLICANT: Marcusson, Eric G.
TITLE OF INVENTION: Antisense Modulation of No. 600
TITLE OF INVENTION: Antisense Modulation of No. 600
TITLE OF INVENTION: Nol-2-Related Proteins
FILE REFERENCE: ISPH-0337
CURRENT APPLICATION NUMBER: US/09/226,568
CURRENT FILING DATE: 1999-01-07
NUMBER OF SEQ ID NOS: 39
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 28
LENGTH: 20
TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
FEATURE: PATENTIAN: Description of Artificial Sequence
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US-09-226-568-28/c
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CURRENT FILING DATE: 1999-01-07
NUMBER OF SEQ ID NOS: 39
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 27
LENGTH: 20
Query Match 0.5%; Score 20; DB 3
Best Local Similarity 100.0%; Pred. No. 9.8;
Matches 20; Conservative 0; Mismatches
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APPLICANT: Bennett, C. Frank
APPLICANT: Dean, Nicholas M.
APPLICANT: Marcusson, Eric G.
TITLE OF INVENTION: Antisense Modulation of
TITLE OF INVENTION: bc1-2-Related Proteins
FILE REFERENCE: ISPH-0337
                                                                                          OTHER INFORMATION: Description of Artificial Sequence: antisense OTHER INFORMATION: sequence
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Sequence 30, Application US/09226568
Patent No. 6001992
GENERAL INFORMATION:
APPLICANT: Ackermann, Elizabeth J.
APPLICANT: Bennett, C. Frank
APPLICANT: Bennett, C. Frank
APPLICANT: Marcusson, Eric G.
TITLE OF INVENTION: Antisense Modulation of No. 6001992el Anti-apoptotic
TITLE OF INVENTION: bcl-2-Related Proteins
FILE REFERENCE: ISPH-0337
CURRENT APPLICATION NUMBER: US/09/226,568
CURRENT FILING DATE: 1999-01-07
NUMBER OF SEQ ID NOS: 39
SOPTWARE: Patentin Ver. 2.0
SEQ ID NO 30
LENGTH: 20
TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: antisense
OTHER INFORMATION: sequence
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APPLICANT: Bennett, C. Frank
APPLICANT: Bennett, C. Frank
APPLICANT: Bennett, C. Frank
APPLICANT: Marcusson, Eric G.
TITLE OF INVENTION: Antisense Modulation of No. 6001992el Anti-apoptotic
TITLE OF INVENTION: bcl-2-Related Proteins
FILE REFERENCE: ISPH-0337
CURRENT APPLICATION NUMBER: US/09/226,568
CURRENT FILING DATE: 1999-01-07
NUMBER OF SEQ ID NOS: 39
SOPTWARE: Patentin Ver. 2.0
SEQ ID NO 29
LENGTH: 20
TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
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US-09-226-568-30/c
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US-09-226-568-29/c
                                                                                    Query Match 0.5%; Score 20; DB 3; Best Local Similarity 100.0%; Pred. No. 9.8; Matches 20; Conservative 0; Mismatches
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OTHER INFORMATION: Description of Artificial Sequence: antisense
OTHER INFORMATION: sequence
                      2231 TGAGTTCCTTCCATTTGACC 2250
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Search completed: March 14, 2003, 08:31:50 Job time: 96.137 secs

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Minimum DB seq length: 0 Maximum DB seq length: 50
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Perfect score:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    OM nucleic - nucleic search, using sw model
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Searched:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Scoring table:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence:
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    OLIGO_NUC Gapext 60.0
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1: //ggn2_6/ptodata/2/pubpna/US07_PUBCOMB.seq:*
2: //ggn2_6/ptodata/2/pubpna/US07_NEW_PUB.seq:*
3: //ggn2_6/ptodata/2/pubpna/US06_PUBCOMB.seq:*
4: //ggn2_6/ptodata/2/pubpna/US06_PUBCOMB.seq:*
5: //ggn2_6/ptodata/2/pubpna/US07_NEW_PUB.seq:*
6: //ggn2_6/ptodata/2/pubpna/US08_NEW_PUB.seq:*
6: //ggn2_6/ptodata/2/pubpna/US08_PUBCOMB.seq:*
7: //ggn2_6/ptodata/2/pubpna/US08_PUBCOMB.seq:*
8: //ggn2_6/ptodata/2/pubpna/US09_PUBCOMB.seq:*
10: //ggn2_6/ptodata/2/pubpna/US09_PUBCOMB.seq:*
11: //ggn2_6/ptodata/2/pubpna/US09_PUBCOMB.seq:*
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12: //ggn2_6/ptodata/2/pubpna/US09_PUBCOMB.seq:*
13: //ggn2_6/ptodata/2/pubpna/US09_PUBCOMB.seq:*
14: //ggn2_6/ptodata/2/pubpna/US09_PUBCOMB.seq:*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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46	46	46	43	34	33	30	30	30	26	25	22	21	21	21	18	17	17	33	Query Match Length DB
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US-10-026-914-10	US-10-026-914-8	US-09-996-634-108	US-10-043-142-4	US-09-811-824-4	US-09-817-513A-22	US-09-954-697-50	US-09-728-466-19	US-09-045-620-3	US-09-811-824-5	US-09-814-986-59	US-10-194-594-12	US-09-828-034-31	US-09-828-034-30	US-09-828-034-11	US-09-263-959-921	US-09-263-959-904	US-09-263-959-900	US-09-884-814-4	ID
	Sequence 8, Appl	Sequence 108, App	Sequence 4,	Sequence 4, A	Sequence 22, Appl	Sequence 50,	Sequence 19, Appl	Sequence 3, Appl	Sequence 5, Appli	Sequence 59, Appl	Sequence 12, Appl	Sequence 31, Appl	Sequence 30,	Sequence 11,	Sequence 921, App	Sequence 904,	Sequence 900	Sequence 4, Appli	Description

44 45	c 43	42	c 41 ·			c 38		c 36	35	c 34	с 33	32	31	30	29	c 28	27			c 24				c 20
14 14	14	14	14	14	14	14	14	14	14	14	14	15	15	15	15	15	15	15	16	16	16	16	16	16
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9 9	9	9	10	10	10	10	10	10	9	10	9	9	9	10	9	9	10	10	10	12	10	9	9	9
US-10-180-762-27 US-09-755-088-1	US-09-804-717A-30	US-09-911-176B-27	US-09-735-995-73	us-09-735-995-17	US-09-263-959-1093	US-09-828-034-28	US-09-828-034-9	US-09-800-629A-27	US-09-863-806-85	US-09-736-863-16	US-09-961-077-1149	US-10-137-316-7	US-10-137-316-5	US-09-007-093-16	US-09-912-263-155	US-09-976-736-27	US-09-873-676-74	US-09-969-373-2606	US-09-935-727-26	US-10-082-018-6	US-09-801-042-8	US-10-026-914-4	US-10-026-914-16	US-10-026-914-14
Sequence 27, Appl Sequence 1, Appli	30,	27,	Sequence 73, Appl	Sequence 17, Appl	Sequence 1093, Ap	Sequence 28, Appl	Sequence 9, Appli	Sequence 27, Appl	Sequence 85, Appl	Sequence 16, Appl	Sequence 1149, Ap	Sequence 7, Appli	Sequence 5, Appli	Sequence 16, Appl	Sequence 155, App	Sequence 27, Appl	Sequence 74, Appl	Sequence 2606, Ap	Sequence 26, Appl	Sequence 6, Appli	Sequence 8, Appli	Sequence 4, Appli	Sequence 16, Appl	Sequence 14, Appl

ALIGNMENTS

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APPLICANT: Chen, Jin-Long
APPLICANT: Chen, Jin-Long
APPLICANT: Amaral, M. Catherine
APPLICANT: Tularik Inc.
TITLE OF INVENTION: Human Uncoupling Protein 2 (hUCP2): Co
TITLE OF INVENTION: Methods of Use
FILE REFERENCE: 018781-0011100S
CURRENT APPLICATION NUMBER: US/09/884,814
CURRENT APPLICATION NUMBER: US/09/884,814
CURRENT FILING DATE: 2001-06-18
PRIOR APPLICATION NUMBER: US 09/124,293
PRIOR FILING DATE: 1998-07-29
NUMBER OF SEQ ID NOS: 8
NUMBER OF SEQ ID NOS: 8
                                                                                                                                                                                                                                                                                                                                                                                                 ; LENGTH: 33
; TYPE: DNA
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Description of Artificial Sequence:U8R primer
US-09-884-814-4
                   Sequence 900, Application US/09263959
Patent NO. US20020150891A1
GENERAL INFORMATION:
APPLICANT: Hood, Leroy E.
APPLICANT: Rowen, Lee
                                                                                                                                  RESULT 2
US-09-263-959-900
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US-09-884-814-4
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; Patent No. US20020127600A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; GENERAL INFORMATION
                                                                                                                                                                                                                                                                                              Query Match 100.0%; Best Local Similarity 100.0%; Matches 17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: PatentIn Ver. 2.1 SEQ ID NO 4
    APPLICANT:
                                                                                                                                                                                                                                                                    453 GGAGCCTCTCGGGAAGC 469
                                                                                                                                                                                                                        17 GGAGCCTCTCGGGAAGC 33
Rowen, Lee
Koop, Ben F.
                                                                                                                                                                                                                                                                                                                                     0.4%; Score 17; DB 10; Length 33; 100.0%; Pred. No. 2e+02;
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                                                                                                                                                                                                                                                                                                                  0
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US-09-263-959-904
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 904, Application US/09263959 Patent No. US20020150891A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ZIP: 98104-7092

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/263,959
FILING DATE: 05-MAR-1999
CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
ATTORNEY/AGENT INFORMATION:
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEFAX: (206) 682-6031
INFORMATION FOR SEQ ID NO: 900:
                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT:
APPLICANT:
                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/263,959
FILLING DATE: 05-MAR-1999
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1148 ACCACCACCACCACCA 1163
ATTORNEY/AGENT INFORMATION:
NAME: MCMASters, David D.
REGISTRATION NUMBER: 33,963
REFERENCE/DOCKET NUMBER: 92
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: ROWEN, LEE
APPLICANT: ROWEN, LEE
APPLICANT: NOTE: DIACHOSTIC AND THERAPEUTIC COMPOSITIONS AND METHODS WHICH UTI
NUMBER OF SEQUENCES: 1279
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: McMasters, David D.
REGISTRATION NUMBER: 33,963
REFERENCE/DOCKET NUMBER: 920010.426C2
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: DI NUMBER OF SEQUENCES: 1 CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                             ZIP:
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                                                                                                                                                                                                                                                                                           COUNTRY:
                                                                                                                                                                                                                                                                                                                  STATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STRANDEDNESS: si
TOPOLOGY: linear
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6300 Columbia Center, 701 Fifth Avenue
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                    920010.426C2
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                                                                                                                                                                                                                                                                                                                                                 701 Fifth Avenue
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0. 6.4e+02;
ches 0;
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RESULT 5
US-09-828-034-11
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US-09-263-959-921
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US-09-263-959-921
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TELEFAX: (206) 682-6031
INFORMATION FOR SEQ ID NO: 904:
SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                          Matches
                                                                                                                                       Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity 100.
Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 921,
Patent No. US
                                                                                                                                                                                                                                                         TELEFAX: (206) 682-603
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
                                                                                          1148 ACCACCACCACCACCA 1163
                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/263,959
FILING DATE: 05-MAR-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1148 ACCACCACCACCACCA 1163
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NAME: McMasters, David D
REGISTRATION NUMBER: 33,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ZIP: 98104-7092
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                        TELEPHONE: (206) 622-4900
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION: DIAGNOSTIC AND THERAPEUTIC COMPOSITIONS AND METHODS WHICH NUMBER OF SEQUENCES: 1279
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                                                                                                                                                                                                                           TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                        REGISTRATION NUMBER: 33,963
REFERENCE/DOCKET NUMBER: 92
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY:
                                                             1 ACCACCACCACCA 16
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6300 Columbia Center, 701 Fifth Avenue
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Koop, Ben F.
                                                                                                                        Conservative
                                                                                                                                                                                                    linear
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100.0%; Pr
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                                                                                                                     Score 16; DB LV; ...; Pred. No. 6.5e+02;
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
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6.4e+02;
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Sequence 11, Application US/09828034 Patent No. US20020064771A1

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RESULT 7
US-09-828-034-31
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US-09-828-034-30
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                                                                                                                                                                                                                                                                                                                                                                                  ; OTHER INFORMATION: Description of Artificial Sequence: Synthetic RNA
US-09-828-034-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION NUMBER: 2001-04-06; PRIOR APPLICATION NUMBER: U.S. 60/195,852; PRIOR FILING DATE: 2000-04-06; NUMBER OF SEQ ID NOS: 33

SOFTWARE: Patentin Ver. 2.1

EQ ID NO 30

LENCTUP A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-09-828-034-11
             GENERAL INFORMATION:
APPLICANT: Zhong, Weldong
APPLICANT: Hong, Zhi
APPLICANT: FERRAL, Eric
TITLE OF INVENTION: HCV REPLICASE COMPLEXES
FILE REFERENCE: INO1165
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 30, Application US/09828034 Patent No. US20020064771A1 GENERAL INFORMATION:
                                                                                                                          Sequence 31, Application US/09828034 Patent No. US20020064771A1
                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity
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Best Local :
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SEQ ID NO 11
LENGTH: 21
                                                                                                                                                                                                                                                                                                               Matches
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APPLICANT: Zhong, Weidong
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Zhong, Weidong
APPLICANT: Hong, Zhi
APPLICANT: Hong, Zhi
APPLICANT: Periari, Eric
TITLE OF INVENTION: HCV REPLICASE COMPLEXES
FILE REFERENCE: INOI165
CURRENT EPPLICATION NUMBER: US/09/828,034
CURRENT FILING DATE: 2001.04-06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION NUMBER: U.S. 60/195,852
PRIOR FILING DATE: 2000-04-06
NUMBER OF SEQ ID NOS: 33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION NUMBER: US/09/828,034
CURRENT FILING DATE: 2001-04-06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Hong, Zhi
APPLICANT: Ferrari, Eric
TITLE OF INVENTION: HCV REPLICASE COMPLEXES
FILE REFERENCE: IN01165
CURRENT APPLICATION NUMBER: US/09/828,034
                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: RNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: RNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LENGTH: 21
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                                                                                                                                                                                                                                                                         1148 ACCACCACCACCACCA 1163
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        OTHER INFORMATION: Description of Artificial Sequence: Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1148 ACCACCACCACCACCA 1163
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local Similarity 100. ses 16; Conservative
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100.0%; Pred. No.
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; Pred. No. 6.6e+02
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. 6.6e+02;
ches 0;
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RESULT 9
US-09-814-986-59
US-09-814-986-59; Sequence 59, Application US/09814986; Sequence 59, Application US/09814986; Patent No. US20020068286A1
GENERAL INFORMATION:
APPLICANT: Kleyn, Patrick W.
MOOTE, Karen J.
                                                                                                                                                                                                                                                                                                                                                                                                                 ; NAME/KBY: misc_feature; LOCATION: (15)...(19); OTHER INFORMATION: Start sequence of the RNA polymerase L protein US-10-194-594-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQ ID NOS: 33
SOFTWARE: PATENTIN VET. 2.1
SEQ ID NO 31
LENGTH: 21
TYPE: RNA
ORGANISM: Artificial Sequence
FEATURE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: US 60/304,125
PRIOR FILING DATE: 2001-07-11
NUMBER OF SEQ ID NOS: 12
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 12
LENGTH: 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 12, Application US/10194594 Publication No. US20030044386A1 GENERAL INFORMATION:
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Best Local :
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Best Local :
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CURRENT APPLICATION NUMBER: US/10/194,594

CURRENT FILING DATE: 2002-07-11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT FILING DATE: 2001-04-06
PRIOR APPLICATION NUMBER: U.S. 60/195,852
PRIOR FILING DATE: 2000-04-06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION: RECOMBINANT VSV FOR THE TREATMENT OF TITLE OF INVENTION: TUMOR CELLS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Barber, Glen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME/KEY: misc_signal LOCATION: (2)...(12) OTHER INFORMATION: Polyadenylation/transcription stop signal of the OTHER INFORMATION: glycoprotein (G) gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: DNA ORGANISM: Vesicular stomatitis virus FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FEATURE:
                                                                                                                                                                                                                                                                                        1866 TTAGTTTTTTTCATAC 1881
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NUMBER OF SEQUENCES: 60
CORRESPONDENCE ADDRESS:
ADDRESSEE: Pennie & Edmonds LLP
                                                          TITLE OF INVENTION: COMPOSITIONS FOR THE TREATMENT AND DIAGNOSIS OF BODY WEIGHT DISORDERS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               16; Conserv
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16; Conserv
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Pred. No.
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Pred. No.
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. 6.6e+02;
ches 0;
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ches 0;
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STREET: 1155 AV CITY: New York STATE: New York

1155 Avenue

of the Americas

COUNTRY: U.S.A.

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APPLICANT: Anderson, George P.

APPLICANT: Mattoussi, Hedi
APPLICANT: Mauro, J. Matthew
APPLICANT: Mauro, J. Matthew
APPLICANT: Sundar, Vikram C.
TITLE OF INVENTION: INORGANIC PARTICLE CONJUGATES
FILE REFERENCE: 01997-282001
CURRENT APPLICATION NUMBER: US/09/811,824
CURRENT FILING DATE: 2001-03-20
PRIOR APPLICATION WHERE: US 60/190,766
PRIOR FILING DATE: 2000-03-20
NUMBER OF SEQ ID NOS: 7
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 5
LENGTH: 26
                                                                                               ; FEATURE:
; OTHER INFORMATION: Synthetically generated primer US-09-811-824-5
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SEQUENCE DESCRIPTION: SEQ ID NO:
US-09-814-986-59
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-09-811-824-5/c
0.4%;
Hest Local Similarity 100.0%;
Matches 16; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 5, Application US/09811824 Publication No. US20020182632A1
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                                                                                                                                                       TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEFAX: (212) 869-97
TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 59:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1148 ACCACCACCACCACCA 1163
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/936,707
APILING DATE: 24-SEP-1997
ATTORNEY/AGENT INFORMATION:
NAME: COTUZZI, LBUITA A.
REGISTRATION NUMBER: 30,742
REFERENCE/DOCKET NUMBER: 7853-100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE CHARACTERISTICS:
LENGTH: 25 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELECOMMUNICATION INFORMATION: TELEPHONE: (212) 790-9090
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US/09/814,986 FILING DATE: 22-Mar-2001 CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (212) 869-9741/8864

 Mismatches

                                 Score 16; DB 9; I
; Pred. No. 6.8e+02;
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                                                     Length 26;
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             Indels
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             0;
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             Gaps
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             0;
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RESULT 13 US-09-954-697-50/c

Sequence 50, Application US/09954697

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; ORGANISM: Artificial Sequence ; FEATURE: THORNMATION: Description of Artificial Sequence: Caspase cDNA ; OTHER INFORMATION: amplification primer US-09-045-620-3
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US-09-728-466-19/c
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US-09-045-620-3/c
Sequence 3, Application US/09045620
Patent No. US20010006793A1
GENERAL INFORMATION:
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                                                                                                                                                                                                                               SOFTWARE: POSEQ ID NO 19
LENGTH: 30
TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: MODULATORS OF EUKARYOTIC CASPASES FILE REFERENCE: 20985.0027/27US
CURRENT APPLICATION UNDERE: U$09/045,620
CURRENT FILING DATE: 2000-03-20
NUMBER OF SEQ ID NOS: 5
SOFTWARE: PATENTIN Ver. 2.1
SEQ ID NO 3
LENGTH: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 19, Application US/09728466 Patent No. US20010029022A1
                                                                    Query Match 0.4
Best Local Similarity 100.
Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
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Best Local Similarity 100.0%; F
Matches 16; Conservative 0;
                                                                                                                                                                                                                                                                                                      FILE REFERENCE: 28341/6216
CURRENT APPLICATION NUMBER: US/09/728,466
CURRENT FILING DATE: 2000-12-01
PRIOR APPLICATION NUMBER: 09/382,616
PRIOR FILING DATE: 1999-08-25
NUMBER OF SEQ ID NOS: 43
COMMANDER: DESCRIPTION NOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: BJORNSTI, Mary-Ann
APPLICANT: HALL, David
APPLICANT: KANG, Jason
                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Fisher, Christopher
APPLICANT: He, Wanxia
TITLE OF INVENTION: Wethods to Identify Anti-Viral Agents
FILE REFERENCE: 28341/6216
                                                                                                                                                                                           ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: DNA
                    1148 ACCACCACCACCACCA 1163
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28 ACCACCACCACCACCA 13
                                                                                                                                                                                                                                                                                           PatentIn Ver. 2.0
                                                                                           100.0%;
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                                                                                           Score 16; DB 10;
; Pred. No. 6.9e+02;
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Pred. No. 6.9e+02;
                                                                             Mismatches
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SEQUENCE CHARACTERISTICS:
LENGTH: 33 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: other nucleic acid
DESCRIPTION: /desc = "primer"
SEQUENCE DESCRIPTION: SEQ ID NO: 22:
US-09-817-513a-22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Allemri, Emad S.
APPLICANT: Allemri, Emad S.
TITLE OF INVENTION: RECOMBINANT, ACTIVE CASPASES AND USSES
TITLE OF INVENTION: THEREOF
FILE REFERENCE: 480140, 431D2
CURRENT PELICATION NUMBER: US/09/954,697
CURRENT FILING DATE: 2001-09-14
NUMBER OF SEQ ID NOS: 116
SOFTWARE: FASTSEQ for Windows Version 3.0
SEQ ID NO 50
SEQ ID NO 50
LENGTH: 30
TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ; FEATURE:
; OTHER INFORMATION: PCR primer for US-09-954-697-50
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    Query Match
Best Local
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                                                                                                                                                                                                                                       NAME: YOUNG J. Suh
REGISTRATION NUMBER: P-41,337
REFERENCE/DOCKET NUMBER: 2786:
TELECOMMUNICATION INFORMATION:
TELEPHONE: 312-474-6300
TELEPHONE: 312-474-0448
INFORMATION FOR SEQ ID NO: 22:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1148 ACCACCACCACCACCA 1163
                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Vers
SOFTWARE: Patentin Release #1.0, Vers
APPLICATION NUMBER: US/09/817,513A
FILING DATE: 28-Jun-2001
CLASSIFICATION: CUNKNOWN>
ATTORNEY/ACENT INFORMATION:
Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  y Match 0.4%; So Local Similarity 100.0%; I hes 16; Conservative 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Lipsky, Brian P.
TITLE OF INVENTION: Cytoplasmic modulators of Integrin
Binding/Signaling
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Staunton, Donald E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COUNTRY: USA
ZIP: 60606
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STATE: Illinois
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CITY: Chicago
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADDRESSEE: Marshall, O'Toc
STREET: 233 South Wacker,
  0.4%;
Score 16; DB 9; I
Pred. No. 6.9e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 16;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           O'Toole, Gerstein, Murray & Borun
cker, 6300 Sears Tower
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6.9e+02;
thes 0;
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                   Length 33;
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Search completed: March 14, 2003, 10:59:24 Job time: 249.857 secs
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                                                                                                                                                                                                     ; OTHER INFORMATION: Synthetically generated primer US-09-811-824-4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 4, Application US/09811824 Publication No. US20020182632A1 GENERAL INFORMATION:
                                                                                                                                    Matches
                                                                                                                                                  Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                       SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Anderson, George P.
APPLICANT: Mattouss1, Hedi
APPLICANT: Mattouss1, Hedi
APPLICANT: Mauro, J. Matthew
APPLICANT: Bawendi, Moungi G.
APPLICANT: Sundar, Vikram C.
TITLE OF INVENTION: INORGANIC PARTICLE CONJUGATES
EILE REFERENCE: 01997-282001
                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION NUMBER: US/09/811,824
CURRENT EILING DATE: 2001-03-20
PRIOR APPLICATION NUMBER: US 60/190,766
PRIOR FILING DATE: 2000-03-20
                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                    LENGTH: 34
TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                1148 ACCACCACCACCACCA 1163
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                                                                   ACCACCACCACCACCA
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                                                                                                                                0.4%; Score 16; nilarity 100.0%; Pred. No. Conservative 0; Mismatci
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Maximum DB
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              OLIGO_NUC Gapop 60.0 , Gapext 60.0
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:/cgn2_6/ptodata/1/pna/US01_COMB.seq:*
:/cgn2_6/ptodata/1/pna/US08_COMB.seq:*
:/cgn2_6/ptodata/1/pna/US091_COMB.seq:*
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:/cgn2_6/ptodata/1/pna/US10B_COMB.seq:*
:/cgn2_6/ptodata/1/pna/US10B_COMB.seq:
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'/cgn2_6/ptodata/1/pna/US6032_COMB.seq *
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/cgn2_6/ptodata/1/pna/US6006_COMB.seq:*
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/cgn2_6/ptodata/1/pna/US6001_COMB.seq:*
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Pred. No. and is greater No. is the number of results predicted by chance to have a greater than or equal to the score of the result being printed, derived by analysis of the total score distribution.

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US-08-472-801-2255	US-08-472-801-2247	US-08-472-801-2231	US-08-668-235-2252	US-08-472-801-2252	US-08-668-235-2242	US-08-668-235-2239	US-08-668-235-2233	US-08-472-801-2242	US-08-472-801-2239	US-08-472-801-2233	US-08-668-235-2262	US-08-668-235-2258	US-08-472-801-2262	US-08-472-801-2258	US-08-668-235-2249	US-08-472-801-2249	US-08-668-235-2235	US-08-472-801-2235	US-08-668-235-2244	US-08-472-801-2244	ID		SUMMARIES
	•	•	Sequence 2252, Ap	Sequence 2252, Ap	Sequence 2242, Ap	Sequence 2239, A	Sequence 2233, A	2242,	•		Sequence 2262, Ap	2258,	Sequence 2262, Ap		Ÿ	2249,	Sequence 2235, Ap		Sequence 2244, Ap	Sequence 2244, Ap	Description		

Gaps

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Sequence 2244, Application US/08668235
GENERAL INFORMATION:
APPLICANT: LATTY J. Smith
TITLE OF INVENTION: Methods and Compositions for
TITLE OF INVENTION: Methods and Compositions for
TITLE OF INVENTION: Reprogramming
FILE REFERENCE: Hesed-1
CURRENT APPLICATION NUMBER: US/08/668, 235
CURRENT FILING DATE: 1996-06-17
EARLIER FILING DATE: 08/23/91
EARLIER FILING DATE: 08/23/91
EARLIER APPLICATION NUMBER: 08/426,781
EARLIER APPLICATION NUMBER: 08/472,801
EARLIER APPLICATION NUMBER: 08/472,801
EARLIER APPLICATION NUMBER: 08/472,801
                                                                                                                                                                                                 US-08-668-235-2244/c
                                                                                                                                                                                                                                                                                                                                                                  ; LENGTH: 46
; TYPE: DNA
; ORGANISM: Homo sapiens
US-08-472-801-2244
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US-08-472-801-2244/c
                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Hesed 2
APPLICANT: Smith, Larry J.
APPLICANT: Smith, Larry J.
TITLE OF INVENTION: Method and Compositions for Cellular
TITLE OF INVENTION: Reprogramming
TILE REFERENCE: Hesed 2
CURRENT APPLICATION NUMBER: US/08/472,801
CURRENT FILING DATE: 1995-06-07
NUMBER OF SEQ ID NOS: 3601
SOFTMARE: FastsEQ for Windows Version 3.0
SEQ ID NO 2244
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 2244, Application US/08472801 GENERAL INFORMATION:
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Best Local Similarity
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ilarity 100.0%; Pred. No. 8.6e-11;
Conservative 0; Mismatches 0;
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US-08-472-801-2253

US-08-472-801-2253

US-08-668-235-2236

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                                                                                                                                                                                                                                                                                                                                    Length 46;
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Sequence 2134, Ap
Sequence 2234, Ap
Sequence 2234, Ap
Sequence 2234, Ap
Sequence 2234, Ap
Sequence 2236, Ap
Sequence 2240, Ap
Sequence 2240, Ap
Sequence 2251, Ap
Sequence 2253, Ap
Sequence 2253, Ap
Sequence 2253, Ap
Sequence 2254, Ap
Sequence 2254, Ap
Sequence 2253, Ap
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Sequence
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ce 2247, Ap
ce 2255, Ap
e 19, Appl
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                                                                  ; TYPE: DNA
; ORGANISM: Homo
US-08-668-235-2235
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; Sequence 2235, Application US/08668235
; GENERAL INFORMATION:
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Sequence 2255, Application US/08472801
SEQUENCE INFORMATION:
APPLICANT: Hesed 2
APPLICANT: Smith, Larry J.
APPLICANT: Smith, Larry J.
TITLE OF INVENTION: Reprogramming
FILE REFERENCE: Hesed 2
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                                                                                                         SOFTWARE: FastSEQ for Windows Version
SEQ ID NO 2235
LENGTH: 42
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION NUMBER: US/08/472,801
CURRENT FILING DATE: 1995-06-07
NUMBER OF SEQ ID NOS: 3601
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 2235
LENGTH: 42
          Best Local Similarity Matches 42; Conserv
                                       Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 1.2
Best Local Similarity 100.
Matches 46; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQ ID NOS: 3629
SOFTWARE: FASTSEQ for Windows Version
SEQ ID NO 2244
LENGTH: 46
                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/08/668,235
CURRENT FILING DATE: 1996-06-17
EARLIER APPLICATION NUMBER: 07/748,997
EARLIER FILING DATE: 08/23/91
EARLIER APPLICATION NUMBER: 08/426,781
EARLIER FILING DATE: 04/22/95
                                                                                                                                                    NUMBER OF SEQ ID NOS:
                                                                                                                                                               EARLIER APPLICATION NUMBER: 08/472,801 EARLIER FILING DATE: 06/07/95
                                                                                                                                                                                                                                                                            APPLICANT: Larry J. Smith
TITLE OF INVERTION: Methods and Compositions for Cellular
TITLE OF INVENTION: Reprogramming
FILE REFERENCE: Hesed-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                 192 CCGGCGAGAGATAGGGGGAGGGGGAGGCCGGCGGCGGTGATTGG 233
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          534 GTCACTACCCTCGACGCCGCCGCCAGCAGAGGAGGAGGAGGAGGACGAG 579
                                                                                                                                                                                                                                                                                                                                                                                                                42 CCGGCGAGAGATAGGGGGAGGGGGAGGCCGGCGCGGTGATTGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           46 GTCACTACCCTCGACGCCGCCGCCAGCAGGAGGAGGAGGACGAG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local
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            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                               sapiens
                         1.1%; Score 42;
100.0%; Pred. No.
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100.0%; Pr
100.0%; Pr
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 42;
Pred. No.
          Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
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                         8e-09;
                                     DB 10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 8;
8e-09;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 10; 1
8.6e-11;
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            0;
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                                    Length 42;
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          Indels
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Gaps

0;

0;

В Q

Matches

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192 CCGGCGAGAGATAGGGGGGAGGGGAGGCCGGCGCGGTGATTGG

233

42 CCGGCGAGAGATAGGGGGAGGGGAGGCCGGCGCGGTGATTGG 1

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CURRENT APPLICATION NUMBER: US/08/668,235
CURRENT FILING DATE: 1996-06-17
EARLIER APPLICATION NUMBER: 07/748,997
EARLIER FILING DATE: 08/23/91
EARLIER FILING DATE: 08/23/91
EARLIER APPLICATION NUMBER: 08/426,781
EARLIER APPLICATION NUMBER: 08/472,801
EARLIER FILING DATE: 06/07/95
NUMBER OF SEQ ID NOS: 3629
SOFTMARE: FASTSEQ FOR WINDOWS Version 3.0
SEQ ID NO 2249
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US-08-668-235-2249/c
; Sequence 2249, Application US/08668235
; GENERAL INFORMATION:
; APPLICANT: Larry J. Smith
; TITLE OF INVENTION: Methods and Compositions for Cellular
; TITLE OF INVENTION: Reprogramming
; FILE REFERENCE: Hesed-1
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RESULT 7
US-08-472-801-2258/c
; Sequence 2258, Application US/08472801
; GENERAL INFORMATION:
; APPLICANT: Hesed 2
; APPLICANT: Smith, Larry J.
; TITLE OF INVENTION: Method and Compositions for Cellular
                                                                                                                                                                                                                                                                                                                                                                     ; TYPE: DNA
; ORGANISM: Homo sapiens
US-08-668-235-2249
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; TYPE: DNA; ORGANISM: Homo sapiens US-08-472-801-2249
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT 5
US-08-472-801-2249/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: Method and Compositions for Cellular TITLE OF INVENTION: Reprogramming FILE REFERENCE: Hesed 2 CURRENT APPLICATION NUMBER: US/08/472,801 CURRENT FILING DATE: 1995-06-07 NUMBER OF SEQ ID NOS: 3601 SOFTMARE: FastSEQ for Windows Version 3.0 SEQ ID NO 2249
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 2249, Application US/08472801 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
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Best Local Similarity
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APPLICANT: Smith, Larry J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH: 40
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                                                                                                                                                                                                                                                                                                                                                                                                                                     ENGTH: 40
                                                                                                                                                                                                                                            3215 GGAAGAACTGCCCTGCCCGCCCATCTCAGAGCCATAAGGT 3254
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     7 Match 1.0%; Score 40; DB 8; L
Local Similarity 100.0%; Pred. No. 7.7e-08;
les 40; Conservative 0; Mismatches 0;
                                                                                                                                                                                                40 GGAAGAACTGCCCTGCCTGCCCATCTCAGAGCCATAAGGT 1
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                                                                                                                                                                                                                                                                                   1.0%;
ilarity 100.0%;
Conservative (
                                                                                                                                                                                                                                                                                   0;
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Pred. No.
                                                                                                                                                                                                                                                                                       Mismatches
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7.7e-08;
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TITLE OF INVENTION: Method and Compositions for Cellular TITLE OF INVENTION: Method and Compositions for Cellular TITLE OF INVENTION: Reprogramming FILE REFERENCE: Hesed 2 FORERIT APPLICATION NUMBER: US/08/472,801 CURRENT FILING DATE: 1995-06-07 NUMBER OF SEQ ID NOS: 3601

SOFTWARE: FastSEQ for Windows Version 3.0

SEQ ID NO 2262

LENGTH: 35

TYPE: DNA

ORGANICOM.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/08/472,801
CURRENT FILING DATE: 1995-06-07
NUMBER OF SEQ ID NOS: 3601
SOFTWARE: FASTSEQ for Windows Version 3.0
SEQ ID NO 2258
SEQ ID NO 2258
LENGTH: 35
TYPE: DNA
ORGANISM: Homo sapiens
US-08-472-801-2258
                                CURRENT APPLICATION NUMBER: US/08/668,235
CURRENT FILING DATE: 1996-06-17
EARLIER APPLICATION NUMBER: 07/748,997
EARLIER FILING DATE: 08/23/91
EARLIER FILING DATE: 08/23/91
EARLIER FILING DATE: 08/426,781
EARLIER FILING DATE: 08/472,801
EARLIER FILING DATE: 06/472,801
EARLIER FILING DATE: 06/07/95
NUMBER OF SEQ ID NOS: 3629
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 2258
LENGTH: 35
TYPE: DATE: 05/07/95
                                                                                                                                                                                                                                                                                                                      US-08-668-235-2258/c
Sequence 2258, Application US/08668235
GEMERAL INFORMATION:
APPLICANT: Larry J. Smith
TITLE OF INVENTION: Methods and Compositions
TITLE OF INVENTION: Reprogramming
FILE REFERENCE: Hesed-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     δÃ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; ORGANISM: Homo sapiens US-08-472-801-2262
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 8
; ORGANISM: Homo sapiens US-08-668-235-2258
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 2262, Application US/08472801 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity
Matches 35; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match 0.9%; Score 35; DB 8; Best Local Similarity 100.0%; Pred. No. 2.2e-(Matches 35; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FILE REFERENCE: Hesed
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         628 GCCACCGGCGCCAAGGACACAAAGCCAATGGGCAG 662
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100.0%; Pred. No. 2.2e-05;
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                                                                                                                                                                                                                                                                                                                                                                     for Cellular
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Query Match Best Local Similarity

Length 35;

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CURRENT APPLICATION NUMBER: US/08/472,801
CURRENT FILING DATE: 1995-06-07
NUMBER OF SEQ ID NOS: 3601
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 2233
LENGTH: 34
TYPE: DNA
ORGANISM: Homo sapiens
US-08-472-801-2233
RESULT 12
US-08-472-801-2239/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: Methods and Compositions
TITLE OF INVENTION: Reprogramming
FILE REFERENCE: Hesed-1
CURRENT APPLICATION NUMBER: US/08/668/235
CURRENT FILING DATE: 1996-06-17
EARLIER APPLICATION NUMBER: 07/48,997
EARLIER FILING DATE: 08/23/91
EARLIER FILING DATE: 04/22/95
EARLIER FILING DATE: 04/22/95
EARLIER FILING DATE: 04/22/95
EARLIER APPLICATION NUMBER: 08/472,801
EARLIER APPLICATION NUMBER: 08/472,801
EARLIER APPLICATION NUMBER: 08/472,801
EARLIER FILING DATE: 06/07/95
NUMBER OF SEQ ID NOS: 3629
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 2262
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; LENGTH: 35
; TYPE: DNA
; ORGANISM: Homo sapiens
US-08-668-235-2262
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         В
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                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 2233, Application US/08472801
GENERAL INFORMATION:
APPLICANT: Hesed 2
APPLICANT: Smith, Larry J.
TITLE OF INVENTION: Method and Compositions for Cellular
TITLE OF INVENTION: Reprogramming
FILE REFERENCE: Hesed 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 2262, Application GENERAL INFORMATION:
APPLICANT: Larry J. Smith
                                                                                                                                                                               Matches
                                                                                                                                                                                                                 Query Match
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                                                                                                  150 CCCGGGAGGGCGACTITTGGCTACGGAGAAGGAG 183
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             628 GCCACCGGCGCCAAGGACACAAAGCCAATGGGCAG 662
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                                                                                                                                                                             Local Similarity 100.0%; ses 34; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           35 GCCACCGGCGCCAAGGACACAAAGCCAATGGGCAG 1
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Similarity 100.0%;
             Application US/08472801
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Application US/08668235
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Methods and Compositions for Cellular
                                                                                                                                                                         0.9%; Score 34; DB 8;
100.0%; Pred. No. 6.9e-(
ive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0;
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 10;
. 2.2e-05;
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                                                                                                                                                                                                               Length 34;
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Sequence 2233, Application US/08668235
GENERAL INFORMATION:
APPLICANT: Larry J. Smith
TITLE OF INVENTION: Methods and Compositions
TITLE OF INVENTION: Reprogramming
FILE REFERENCE: Hesed-1
CURRENT APPLICATION NUMBER: US/08/668,235
CURRENT FILING DATE: 1996-06-17
EARLIER APPLICATION NUMBER: 07/748,997
EARLIER FILING DATE: 08/23/91
EARLIER FILING DATE: 08/23/91
EARLIER APPLICATION NUMBER: 08/426,781
EARLIER APPLICATION NUMBER: 08/472,801
EARLIER APPLICATION NUMBER: 08/472,801
EARLIER FILING DATE: 06/07/95
NUMBER OF SEQ ID NOS: 3629
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 2333
LENGTH: 34
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; ORGANISM: Homo sapiens US-08-472-801-2242
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SCOTTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 2239
LENGTH: 34
TYPE: DNA
CRGANTSM: Homo sapiens
US-08-472-801-2239
                                                                                                                                                                                                                                                                                                                                                                                 US-08-668-235-2233/c
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US-08-472-801-2242/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION NUMBER: US/08/472,801
CURRENT FILING DATE: 1995-06-07
NUMBER OF SEQ ID NOS: 3601
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 2242
LENTH: 34
TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 2242, Application US/08472801
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Hesed 2

APPLICANT: Smith, Larry J.

APPLICANT: Smith, Larry J.

TITLE OF INVENTION: Method and Compositions for Cellular

TITLE OF INVENTION: Reprogramming

FILE REFERENCE: Hesed 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Hesed 2
APPLICANT: Smith, Larry J.
TITLE OF INVENTION: Method and Compositions
TITLE OF INVENTION: Reprogramming
FILE REFERENCE: Hesed 2
CURRENT APPLICATION NUMBER: US/08/472,801
CURRENT FILING DATE: 1995-06-07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         485 CGCTGCTGGAGTTGGTCGGGGAATCTGGTAATAA 518
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         260 CCACCCTCACGCCAGACTCCCGGAGGGTCGCGCG 293
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 34 CGCTGCTGGAGTTGGTCGGGGAATCTGGTAATAA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local
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Similarity 100.0%; Pred. No. o...
? ? ? ? ? Mismatches
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Conservative
                                                                                                                                                                                                                                                                                            Methods and Compositions for Cellular
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Pred. No. 6.9e-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 34;
Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   6.9e-05;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 34;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0,:
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RESULT 15

US-08-668-235-2239/c

Sequence 2239, Application US/08668235

GENERAL INFORMATION:

APPLICANT: LALTY J. Smith

TITLE OF INVENTION: Methods and Compositions for Cellular

TITLE OF INVENTION: Methods and Compositions for Cellular

TITLE OF INVENTION: Methods and Compositions for Cellular

CURRENT APPLICATION NUMBER: US/08/668,235

CURRENT FILING DATE: 1996-06-17

EARLIER APPLICATION NUMBER: 07/748,997

EARLIER APPLICATION NUMBER: 08/426,781

EARLIER FILING DATE: 08/23/91

EARLIER FILING DATE: 08/23/91

EARLIER FILING DATE: 08/07/95

UNMBER OF SEQ ID NOS: 3629

SOPTWARE: FastSEQ for Windows Version 3.0

LENGTH: 34

TYPE: DNA

ORGANISM: Homo sapiens
Search completed: March 14, 2003, 10:28:01 Job time: 5797.01 secs
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; TYPE: DNA
; ORGANISM: Homo sapiens
US-08-668-235-2233
                                                                                                                                                                             Query Match 0.9%; Score 34; DB 10; Length 34; Best Local Similarity 100.0%; Pred. No. 6.9e-05; Matches 34; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 0.9%; Score 34; DB 10; Length 34; Best Local Similarity 100.0%; Pred. No. 6.9e-05; Matches 34; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              150 CCCGGGAGGGCGACTTTTGGCTACGGAGAAGGAG 183
                                                                                       0;
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Title: Perfect score: Sequence:

US-09-869-894-18 3934 1 tccagtaaggagtc

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nucleic search, using sw

model

GenCore version 5.1.4_p5_4578 Copyright (c) 1993.- 2003 Compugen Ltd.

Scoring table:

OLIGO_NUC Gapop 60.0 , Gapext 60.0

4666902 seqs, 1136064160 residues

Total number of hits satisfying

chosen

parameters:

Word size Searched:

Minimum DB Maximum DB

seq

length:

50

Post-processing: Listing first 45 summaries

Pending_Patents_NA_New: *

987654321

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 /cgn2_6/ptodata/2/pna/US07_NEW_COMB.seq:*
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 /cgn2_6/ptodata/2/pna/US10_NEW_COMB.seq:*
 /cgn2_6/ptodata/2/pna/US10_NEW_COMB.seq:*

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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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                        2555, App
27749, A
254577,
264577,
27862, Ap
27862, Ap
27862, Ap
27862, Ap
27862, Ap
27862, Appl
27862
APPLICANT: Shimkets, Richard A
APPLICANT: Smithson, Glennda
TITLE OF INVENTION: THERAPEUTIC POLYPEPTIDES, NUCLEIC ACI
FILE REFERENCE: 21402-442A
CURRENT APPLICATION NUMBER: US/10/236,392
CURRENT FILING DATE: 2002-09-06
PRIOR APPLICATION NUMBER: US/9/540,763
PRIOR FILING DATE: 2000-03-30
PRIOR FILING DATE: 2000-03-30
PRIOR APPLICATION NUMBER: US/9/390,155
PRIOR APPLICATION NUMBER: US/9/635,949
PRIOR APPLICATION NUMBER: US/9/635,949
PRIOR APPLICATION NUMBER: US/9/635,949
PRIOR FILING DATE: 2000-08-10
PRIOR FILING DATE: 2000-09-12
PRIOR APPLICATION NUMBER: US/9/318,765
PRIOR FILING DATE: 2001-09-12
PRIOR APPLICATION NUMBER: US/9/357,303
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US-10-236-392-256/c
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APPLICANT: Boldog, Fe
APPLICANT: Burgess, C
APPLICANT: Casman, St
APPLICANT: Casman, St
APPLICANT: Cabtree,
APPLICANT: Crabtree,
APPLICANT: Edinger,
APPLICANT: Edinger,
APPLICANT: Ellerman,
APPLICANT: Gorman, Li
APPLICANT: Gorman, Li
APPLICANT: Gusev, Vla
APPLICANT: Gusev, Vla
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APPLICANT:
APPLICANT:
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Rastelli, Luca
Reiger, Daniel K
Rothenberg, Mark E
Shenoy, Suresh
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gorman, Linda
Grosse, William M
Gusev, Vladamir
                                                                                                                                                                                                                                                                                                                                                                                                                                             Miller, Charles |
Millet, Isabelle
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Ellerman, Karen
Gerlach, Valerie
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Edinger, Shlomit,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Chapoval, Andrei
                                                                                                                                                                                                                                                                                                                                                                                         Patturajan, Meera
Pena, Carol A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MacDougall, John
Malyankar, Uriel
Miller, Charles H
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Catterton, Elina
                                                                                                                                                                                                                                                                                                                                                                                                                               Padigaru, Muralidhara
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LaRochelle, William
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Kekuda, Ramesh
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Boldog, Ferenc L
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Catherine,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   H
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US-60-449-155-609
US-10-303-778-1689
US-10-310-188-31362
US-10-310-188-79759
US-10-310-188-48654
US-10-310-188-62082
US-10-310-188-62082
US-10-310-188-62082
US-10-355-577-7439
US-10-355-577-476248
US-10-355-577-870929
US-10-355-577-808-15971
US-60-427-808-179914
US-60-427-808-179914
US-60-427-808-179914
US-60-427-808-321372
US-10-287-787-13609
US-10-050-902-299
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8706248,
870629,
136571,
179014,
409599,
65488,
321372,
33609,
297, App
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609, App
16089, A
31362, A
255, App
79759, A
486759, A
62082, A
70435, Ap
                                                                                                                                                                                                                                  SAME
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Result No.

Query Match

Length

DB

SUMMARIES

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0 0 0 0 0 0 0 0

0.0

US-10-236-392-256
US-10-233-38-8053
US-10-310-188-79749
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US-10-310-188-7862
US-10-310-188-7969
US-10-310-188-79447
US-10-310-188-81944
US-10-19-93-38-357
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US-10-310-188-79750
US-10-310-188-79750
US-10-310-188-79750
US-60-427-808-537621

Sequence Sequence Sequence Sequence

Sequence Sequence Sequence Sequence Sequence

Description

Sequence

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RESULT 3
US-10-310-188-79749/c
US-10-310-188-79749/c
Sequence 79749, Application US/10310188
GENERAL INFORMATION:
APPLICANT: RosettaGemonics
APPLICANT: ROSettaGemonics
TITLE OF INVENTION: BIOINFORMATICALLY DETECTABLE GROUP OF NOVEL VIRAL REGULATORY GENE
TITLE OF INVENTION: USES THEREOF
FILE REFERENCE: 47487
FILE REFERENCE: 47487
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 2
US-10-293-338-8053/c;
Sequence 8053, App.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR FILING DATE: 2002-02-15
PRIOR APPLICATION NUMBER: US60/367,753
PRIOR APPLICATION NUMBER: US60/369,479
PRIOR APPLICATION NUMBER: US60/369,479
PRIOR APPLICATION NUMBER: US60/369,479
PRIOR PILING DATE: 2002-04-02
PRIOR FILING DATE: 2000-09-12
PRIOR FILING DATE: 2000-09-12
PRIOR APPLICATION NUMBER: US60/318,120
PRIOR APPLICATION NUMBER: US60/318,130
PRIOR APPLICATION NUMBER: US60/318,130
PRIOR APPLICATION NUMBER: US60/318,130
PRIOR FILING DATE: 2001-09-07
PRIOR FILING DATE: 2001-09-07
Remaining Prior Application data removed - Securitable Of SEQ ID NOS: 794
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Qy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; TYPE: DNA; Homo sapiens; ORGANISM: Homo sapiens US-10-293-338-8053
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GENERAL INFORMATION:
APPLICANT: ROSEITAGENOMICS LTD
TITLE OF INVENTION: BIOINFORMATICALLY DETECTABLE GROUP OF NOVEL REGULATORY GENES AND
TITLE OF INVENTION: THEREOF
FILE REFERENCE: 45282
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION NUMBER: US/10/293,338
CURRENT FILING DATE: 2002-11-14
NUMBER OF SEQ ID NOS: 8785
SOFTWARE: PatentIn version 3.1
SEQ ID NO 8053
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: Custom
SEQ ID NO 256
LENGTH: 25
NUMBER OF SEQ ID NOS: 86841
SOFTWARE: Patentin version 3.1
SEQ ID NO 79749
LENGTH: 24
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Best Local S
                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
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                                                                                        CURRENT APPLICATION NUMBER: US/10/310,188 CURRENT FILING DATE: 2002-12-19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LENGTH: 24
                                                                                                                                                                                                                                                                                                                                                                                            1330 AGTTTCAGCAACAAACAAACTTTG 1353
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                                                                                                                                                                                                                                                                                                                                                    24 AGTTTCAGCAACAAACAAACTTTG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                   y Match 0.6%;
Local Similarity 100.0%;
nes 24; Conservative (
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les 25; Conserv
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0.079;
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CURRENT APPLICATION NUMBER: US/10/310,188
CURRENT FILING DATE: 2002-12-19
NUMBER OF SEQ ID NOS: 86841
SOFTWARE: PatentIn version 3.1
SEQ ID NO 2068
LENGTH: 22
TYPE: DNA
CORGANISM: Homo sapiens
US-10-310-188-2068
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; SOFTWARE: Microarray Probe Sequence Listing
; SEQ ID NO 454577
LENGTH: 25
TYPE: DNA
; ORGANISM: Rattus norvegicus
US-60-427-836-454577
RESULT 6
US-10-310-188-7862/c
Sequence 7862, Application US/10310188
; GENERAL INFORMATION:
APPLICANT: RosettaGemonics
, APPLICANT: ROSETION: BIOINFORMATICALLY DETECTABLE GROUP OF NOVEL VIRAL REGULATORY G
, TITLE OF INVENTION: USES THEREOF
; FILE REFERENCE: 47487
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US-10-310-188-2068/c
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Best Local S
Matches 23
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GENERAL INFORMATION:
APPLICANT: RosettaGemonics
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Best Local Similarity 100.0%;
Matches 24; Conservative
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Best Local Similarity
Matches 22; Conserva
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TITLE OF INVENTION: Methods of Genetic Analysis of Rat
FILE REFERENCE: 3527
CURRENT APPLICATION NUMBER: US/60/427,836
CURRENT FILING DATE: 2002-11-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: BIOINFORMATICALLY DETECTABLE GROUP OF NOVEL VIRAL REGULATORY TITLE OF INVENTION: USES THEREOF FILE REFERENCE: 47487
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Local Similarity 100.0%; Pred. No.
Les 23; Conservative 0; Mismato
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100.0%; Pred. No.
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CURRENT APPLICATION NUMBER: US/10/355,577
CURRENT FILING DATE: 2003-01-31
NUMBER OF SEQ ID NOS: 997516
SOFTWARE: Microarray Probe Sequence Listing Generator V 1.1
SEQ ID NO 900097
LENGTH: 25
TYPE: DNA
CORGANIZM: Homo sapien
US-10-355-577-900097
                                                                                                                                                                                                                                                                                        US-10-355-577-900097
Sequence 900097, Application US/10355577
GENERAL INFORMATION:
APPLICANT: Mittmann, Michael
TITLE OF INVENTION: Methods of Genetic Analysis of
FILE REFERENCE: 3121
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; ORGANISM: Homo sapiens
US-10-310-188-73447
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US-10-310-188-73447/c
US-10-310-188-73447, Application US/10310188
GENERAL INFORMATION:
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RESULT 9
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CURRENT FILING DATE: 2002-12-19
NUMBER OF SEQ ID NOS: 86841
SOFTWARE: Patentin version 3.1
SEQ ID NO 7862
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Matches
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Best Local :
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CURRENT FILING DATE: 2002-12-19
NUMBER OF SEQ ID NOS: 86841
SOFTWARE: Patentin version 3.1
SEQ ID NO 73447
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Best Local :
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Best Local
                                                                                                    Matches
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TITLE OF INVENTION: USES THEREOF
FILE REFERENCE: 47487
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Sequence 57, Application US/10169983
GENERAL INFORMATION:
APPLICANT: Takara Shuzo Co., Ltd.
TITLE OF INVENTION: Therapeutic agents
FILE REFERENCE: 01-011-PCT
CURRENT APPLICATION UNMER: US/10/169,983
CURRENT FILING DATE: 2002-07-14
PRIOR APPLICATION NUMBER: UP 2000-4989
PRIOR FILING DATE: 2000-01-13
PRIOR APPLICATION NUMBER: JP 2000-303711
PRIOR APPLICATION NUMBER: JP 2000-303711
PRIOR APPLICATION NUMBER: JP 2000-303711
PRIOR FILING DATE: 2000-10-03
NUMBER OF SEQ ID NOS: 61
SEQ ID NO 57
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US-10-310-188-81944/c
US-10-310-188-81944/c
Sequence 81944, Application US/10310188
GENERAL INFORMATION:
APPLICANT: RosettaGemonics
TITLE OF INVENTION: BIOINFORMATICALLY DETECTABLE GROUP OF NOVEL VIRAL REGULATORY
TITLE OF INVENTION: USES THEREOF
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US-10-169-983-57
; Sequence 57, A
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// ORGANISM: Homo sapiens
US-10-310-188-81944
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CURRENT APPLICATION NUMBER: US/10/310,188
CURRENT FILING DATE: 2002-12-19
NUMBER OF SEQ ID NOS: 86841
SOFTWARE: PATENTIN VERSION 3.1
SEQ ID NO 81944
LENGTH: 20
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CURRENT FILING DATE: 2002-11-14
NUMBER OF SEQ ID NOS: 8785
SOFTWARE: PATENTIN VERSION 3.1
SEQ ID NO 3254
LENGTH: 20
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APPLICANT: RosettaGenomics LTD
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FEATURE: OTHER INFORMATION: Designed primer based on nucleotide sequence of
                                      TYPE: DNA
ORGANISM: Artificial Sequence
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                                                                                      ENGTH: 20
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Local Similarity 100.0%;
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; OTHER INFORMATION: human myeloid leukemia cell differentiation US-10-169-983-57

8;

Length 20;

protein-1

mRNA.

Query Match

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GENERAL INFORMATION:

APPLICANT: Xue Mei Zhou

TITLE OF INVENTION, Methods of Genetic Analysis of Mouse

FILE REFERENCE: 3528

CURRENT APPLICATION NUMBER: US/60/427,808

CURRENT APPLICATION NUMBER: US/60/427,808

CURRENT FILING DATE: 2002-11-20

NUMBER OF SEQ ID NOS: 982914

SOFTWARE: Microarray Probe Sequence Listing Generator V 1.1

SEQ ID NO 23817

LENGTH: 25

TYPE: DNA

ORGANISM: Mus musculus

US-60-427-808-23817
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 58, Application US/10169983
GENERAL INFORMATION:
APPLICANT: Takara Shuzo Co., Ltd.
TITLE OF INVENTION: Therapeutic agents
FILE REFERENCE: 01-011-PCT
CURRENT APPLICATION NUMBER: US/10/169,983
CURRENT FILING DATE: 2002-07-14
PRIOR APPLICATION NUMBER: JP 2000-4989
PRIOR FILING DATE: 2000-01-13
PRIOR FILING DATE: 2000-01-03
INUMBER OF SEQ ID NOMBER: JP 2000-303711
PRIOR FILING DATE: 2000-10-03
SEQ ID NO 58
LENGTH: 20
TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE: TURNORMATTON. Footland Services
RESULT 14
US-10-293-338-1275/c
; Sequence 1275, Application US/10293338
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US-60-427-808-23817; Sequence 23817, AJ
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CURRENT APPLICATION NUMBER: US/10/310,188
CURRENT FILING DATE: 2002-12-19
NUMBER OF SEQ ID NOS: 86841
SOFTWARE: PatentIn version 3.1
SEQ ID NO 57069
LENGTH: 19
TYPE: DNA
ORGANISM: Homo Sapiens
US-10-310-188-57069
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TITLE OF INVENTION: BIOINFORMATICALLY DETECTABLE GROUP OF NOVEL REGULATORY GENES
TITLE OF INVENTION: THEREOF
FILL REFERENCE: 45282
CURRENT APPLICATION NUMBER: US/10/293,338
CURRENT FILING DATE: 2002-11-14
NUMBER OF SEQ ID NOS: 8785
SOFTWARE: PatentIn version 3.1
SEQ ID NO 1275
LENGTH: 19
TYPE: DNA
ORGANISM: Homo sapiens
US-10-293-338-1275
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US-10-310-188-57069/, Application US/10310188
; Sequence 57069, Application US/10310188
; GENERAL INFORMATION:
; APPLICANT: ROSETTAGEMONICS
; TITLE OF INVENTION: BIGINFORMATICALLY DETECTABLE GROUP OF NOVEL VIRAL REGULATORY G
; TITLE OF INVENTION: USES THEREOF
; FILE REFERENCE: 47487
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Search completed: March 14, 2003, 10:49:54
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Maximum DB seq length: 50
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AZ365602 lM0112017
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AZ304663 1M0004016
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ALIGNMENTS

AA463890

DEFINITION aa08003.51 Soares_NhHMPu_S1 Home sapiens cDNA clone IMAGE:812645 3'

DEFINITION aa08003.51 Soares_NhHMPu_S1 Home sapiens cDNA clone IMAGE:812645 3'

SINCL HUMAN Q07820 INDUCED WYELOID LBUKEMIA CELL

ACCESSION AA463890.1 GI:2188774

KEYWORDS

SOURCE

ORGANISM Edwaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

Mammalla; Butheria; Primates; Catarrhin; Hominidae; Homo.

REFERENCE 1 (bases 1 to 31)

AUTHORS

KUCABA,T., Lacy,M., Lennon,G., Marra,M., Martin,J., Moore,B., Schellenberg,K., Steptoe,M., Tan,F., Thelsing,B., White,Y., Rylie

TITLE

JOURNAL

Contact: Wilson R.

Washington University School of Medicine

4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108

Tel: 314 286 1810

Fax: 314 286 1810

Fmail: estewatson.wustl.edu

This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further information.

Possible reversed clone: similarity on wrong strand

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JOURNAL COMMENT
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AUTHORS
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KEYWORDS
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Rukaryota; Metazoa; Chordata; Craniata; verturalizate; Muridae; Murinae; Murinae; Rodentia; Sciurognathi; Muridae; Murinae; Murinae; Murinae; Rodentia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Murinae; Rodentia; Chases 1 to 26; 1 (bases 1 to 26; 1)
2 (bases 1 to 26; 1)
3 (bases 1 to 26; 1)
4 (bases 1 to 26; 1)
5 (bases 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    plasmid inserts
Unpublished (2000)
Contact: Robert B. Weiss
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High quality sequence stop: 19.
Location/Qualifiers
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                                                                                                                        High quality sequence stop: 26.
Location/Qualifiers
                                                                                                                                                                                         Email: ddunn@genetics.utah.edu
Insert Length: 10000 std Error:
Plate: 0112 row: 0 column: 17
Seq primer: CGTTGTAAAACGACGGCCAGT
Class: plasmid ends
                                                                                                                                                                                                                                                                                                                                                    Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                       University of Utah Genome Center
University of Utah
Rm. 308, Biomedical Polymers Res
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                                                                                                                                                                                                                                                                                                                                                                                                               USA
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/organism="Mus musculus"
/strain="C57BL/6J"
/db_xref="taxon:10090"
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/tissue_type="Pooled human melanocyte, fetal heart, and
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/clone="IMAGE:812645"
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/db_xref="GDB:6043555"
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Best Local Similarity
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1M0076C22R Mouse 10kb plasmid UUGC1M library Mus musculus genomic clone UUGC1M0076C22 R, DNA sequence.
A2343185
                                                                                                                                                                                      Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0076 row: C column: 22
Seq primer: CACACAGGAAACAGCTATGACC
                                                                                                                                                                                                                                                                                                                                                                                                          Contact: Robert B. University of Utah University of Utah
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                                                                                                                                   High quality sequence stop: 34.
                                                                                                                                                                                                                                                                                                    Tel: 801 585 5606
Fax: 801 585 7177
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                                                                                                                                                                 plasmid ends
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/clone_lib="Mouse 10kb plasmid UUGC1M library"
/sex="Male"
/strain="C57BL/6J"
/db_xref="taxon:10090"
                                                                                                        Location/Qualifiers
                                                     /organism="Mus musculus"
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Genome Center
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Pred. No.
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UUGC1M library Mus musculus genomic

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Pedersen, T., Reilly

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                                                                                                                                                                                                                                                                                                                                                                                                                   Email: cgapbs-r@mail.nih.gov
Life Technologies catalog #: 11548-013
DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                                                                                                                                                      Tumor Gene Index
Unpublished (1997)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 (bases 1 to 49)
NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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AI569150
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                                /note="Organ: pancreas; Vector: pCMV-SPORT6; Site_1: SalI Site_2: NotI; Cloned unidirectionally. Primer: Oligo dT. Average insert size 1.72 kb. Life Technologies catalog #: 11548-013"

7 c 2 g 22 t
                                                                                                                                                             /clone_lib="NCI_CGAP_Pan1"
/tissue_type="adenocarcinoma"
/lab_host="DH10B"
                                                                                                                                                                                                                                            /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="IMAGE:2224876"
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/note="Vector: PWD42nv; Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jackson
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/clone_lib="Mouse 10kb plasmid UUGC1M library"
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97;
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Best Local Similarity 100.0%; Pred. No.
Matches 19; Conservative 0; Mismatcl
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Unpublished (2000)
Contact: Robert B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0004 row: 0 column: 16
Seg primer: CACACAGGAAACAGCTATGACC
Class: plasmid ends
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University of Utah
Rm. 308, Biomedical
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Location/Qualifiers
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Fax: 801 585 7177
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(http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with 74 DNA polymerase and 74 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gil4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid Rl. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to chemically-competent E. coli XLIO-Gold (Stratagene) cells and selected for ampicillin resistance."
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /organism="Mus musculus"
/strain="C57BL/6J"
/db_xref="taxon:10090"
/clone="UUGC1M0004016"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /clone_lib="Mouse 10kb plasmid UUGCIM library"
/sex="Male"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    'lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"
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SOURCE

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DEFINITION

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BASE COUNT

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        BASE COUNT
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Best Local Similarity
Matches 18; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TITLE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               44 bp DNA linear GSS 04-OCT-2011M0254P10R Mouse 10kb plasmid UUGC1M library Mus musculus genomic clone UUGC1M0254P10 R, DNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
Rm. 308, Biomedical Polymers Research Bldg., 20 S. 2030 E., SLC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GSS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error:
Plate: 0254 row: P column: 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mus musculus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Seq primer: CACACAGGAAACAGCTATGACC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            plasmid inserts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    and Wright, D., Weiss, R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                house mouse.
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801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        quality sequence stop: 44.
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(http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gil4732114|gbhAF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /strain="C57BL/6J"
/db_xref="taxon:10090"
                                                                                                                                                                                                                                                                                                                                                                                                                                            /sex="Male"
                                                                                                                                                                                                                                                                                                                                                                                                                                                             /clone_lib="Mouse 10kb plasmid UUGC1M library"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /clone="UUGC1M0254P10"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /organism="Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Location/Qualifiers
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Pred. No.
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3.1e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 plasmid inserts
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Mammalia; Eutheria;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Class: plasmid ends
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Tel: 801 585 5606
Fax: 801 585 7177
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                                                                                                                                                                                                                                                                                                                                                                                                         /note="Vector:
                                                                                                                                                                                                                                                                                                                                                                                                                                                /sex="Male"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GI:10518157
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ORIGIN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Email: ddunnigenetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0170 row: P column: 09
Seq primer: CACACAGGAAACAGCTATGACC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          I (bases I to 28)

1 (bases I to 28)

Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Rei Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Rei M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, M., Rose, M., Rose, R., Stokes, R.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         High quality sequence stop: 28.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
Rm. 308, Biomedical Polymers Research Bldg.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mouse whole genome scaffolding with paired end reads
                                                                                                                            (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWDA2 (gi14732114(gb)AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and the charged adaptored mouse DNA was annealed to
purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
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/clone="UUGC1M0170P09"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /organism="Mus musculus"
/strain="C57BL/6J"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /clone_lib="Mouse 10kb plasmid UUGC1M library"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"
/note="Vector: PWD42nv; Purified genomic DNA from M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Sutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 18; DB 17; L; Pred. No. 3.1e+03;
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on Niederhausern,A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Insert Size: 953
High quality sequence starts: 1
High quality sequence stops: 1
Source: IMAGE Consortium, LLNL
This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further information.
Trace considered overall poor quality
Trace considered clone: Similarity on wrong strand
Insert Length: 953
Std Error: 0.00
Seq primer: M13Rev
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Hillier, L., Clark, N., Dubuque, T., Elliston, K., Hawkins, M., Holman, M., Kucaba, T., Le, M., Lennon, G., Marra, M., Parsons, Rifkin, L., Rohlfing, T., Soares, M., Tan, F., Trevaskis, E., Waterst, R., Williamson, A., Wohldmann, P. and Wilson, R.

The WashU-Merck EST Project
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Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Tel: 314 286 1800
Fax: 314 286 1810
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Washington University School of Medicine
4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    EST
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Email: est@watson.wustl.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Contact: Wilson RK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Unpublished (1995)
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1 (bases 1 to 40)
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                                                                                                                                                                     /dev_stage="adult"
//lab_host="DH10B (ampicillin resistant)"
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//lab_host="DH10B (ampicillin resistant)"
//lab_host="DH10B (ampicillin resistant)
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/clone="IMAGE:161913"
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Eutheria; Primates; Catarrhini; Hominidae; Homo
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Pred. No.
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Insert Length: 10000 Std Error:
Plate: 046 row F column: 19
Seq primer: CGTTGTAAAACGACGGCCAGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.
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Fax: 801 585 7177
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Contact: Robert B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            University of Utah Genome Center University of Utah
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                                                                                                                                                                                              adaptored vector DNA, and transformed into chemically-competent E. coli XLIO-Gold (Stratagene) cells and selected for ampicillin resistance."

2 c 12 g 22 t
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /db_xref="taxon:10090"
/clone="UUGCIM0496F19"
/clone_lib="Mouse 10kb plasmid UUGC1M library"
/sex="Male"
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/strain="C57BL/6J"
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AZ315293/c
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Query Match
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AZ315293
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D.,Weiss,R.
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University of Utah
Rm. 308, Biomedical Polymers Reso
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High quality sequence stop: 19.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error:
Plate: 0032 row: p column: 20
Seq primer: CGTTGTAAAACGACGGCCAGT
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Fax: 801 585 7177
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                                                                                                                                                                                 (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWDA2 (gil4732114[gb]AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and
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                                                                            purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XLIO-Gold (Stratagene) cells and selected for ampicillin resistance."
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             /note-"Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson
                                                                                                                                                                                                                                                                                                                                                                                                                                           Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /db_xref="taxon:10090"
/clone="UUGC1M0032P20"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /organism="Mus musculus"
/strain="C57BL/6J"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /lab_host="E. Coli strain XL10-Gold, T1-resistant,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /sex="Male"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /clone_lib="Mouse 10kb plasmid UUGC1M library"
0.4%;
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DB 17; Length 19; 3.4e+04;
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Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
Oniversity of Utah
Rm. 308, Biomedical Polymers Rese
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Fax: 801 585 7177
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Insert Length: 10000 Std Error: 0.00
Plate: 0455 row: A column: 24
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Mammalia; Eutheria; Rodentia;
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                                                                                                                                                                                                     purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."

1 0 c 14 g 6 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                            10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pmD42 (gil4732114|gb|AF129072-1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to appropriate the constant of the co
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/strain="C57BL/6J"
/db_xref="taxon:10090"
/clone="UUGC1M0455A24"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Laboratory Mouse DNA Resource
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (http://www.jax.org/resources/documents/dnares/). The DNA
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DB 17; Length 20; 3.4e+04;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: (
Plate: 0027 row: P column: 11
Seq primer: CACACAGGAAACAGCTATGACC
Class: plasmid ends
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     High quality sequence stop: 23.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Unpublished (2000)
Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
Rm. 308, Biomedical Polymers Reso
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D.,Weiss,R.
Mouse whole genome scaffolding with paired end reads from 10kb plasmid inserts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Tel: 801 585 5606
Fax: 801 585 7177
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                                                                                          10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gil4732114(gb)AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
                                                                                                                                                                                                                                                                                                                                                (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"
/note="Vector: PWD42nv; Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jackson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Biomedical Polymers Research Bldg., 20 S. 2030 E., SLC, UT
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             0.4%;
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          DB 17;
. 3.3e+04;
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Insert Length: 10000 Std Error:
Plate: 0167 row: A column: 16
Seq primer: CGTTGTAAAACGACGGCCAGT
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1 (bases 1 to 23)
Dunn, D., Aoyaqi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.
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University of Utah
Rm. 308, Biomedical Polymers Research Bldg.,
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Contact: Robert B.
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Fax: 801 585 7177
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2M0167A16F Mouse 10kb plasmid UUGC1M
clone UUGC2M0167A16 F, DNA sequence.
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AZ860972.1 GI:13056826
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quality sequence stop: 23.
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                                                                                          adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gi|4/32114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
                                                                                                                                                                                                                                                                                                                                                          /note="Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson Laboratory Mouse DNA Resource (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The
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/clone="UUGC2M0167A16"
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/strain="C57BL/6J"
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0.4%;
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DB 17; Length 23; 3.3e+04;
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RESULT 14
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Best Local Similarity
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University of Utah
Rm. 308, Biomedical Polymers Res
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Unpublished (2000)
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clone UUGC1M0248M02 F, DNA sequence.
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Location/Qualifiers
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Insert Length: 10000 Std Error:
Plate: 0248 row: M column: 02
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Mammalla; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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Fax: 801 585 7177
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                                                                                           chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance.  
 15 c 1 g 1 t
                                                                                                                                                                  10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pMD42 gil4732114|pb|RF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"
/note="Vector: PWD42nv; Purified genomic DNA from M.
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/strain="C57BL/6J"
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Mouse whole genome scaffolding with paired end reads from 10kb
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AZ405428
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Unpublished (2000)
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Insert Length: 10000 Std Error: 0.00
Plate: 0174 row: D column: 13
Seq primer: CGTTGTAAAACGACGGCCAGT
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University of Utah
Rm. 308, Biomedical Polymers Res
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus. 1 (bases 1 to 26)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GSS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          High quality sequence stop: 26.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Contact: Robert B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AZ405428.1 GI:10529441
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Class: plasmid ends
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        house mouse
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                                                                                                                          (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWDA2 (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and preparative plasmid R1.
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adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicullin resistance."

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                                                                                                   purified. The sheared, adaptored mouse DNA was annealed to
                                                                                                                                                                                                                                                                                                                                                                                                                                                         /note="Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Biomedical Polymers Research Bldg., 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /organism="Mus musculus"
/strain="C57BL/6J"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /clone_lib="Mouse 10kb plasmid UUGC1M library"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /db_xref="taxon:10090"
/clone="UUGC1M0174D13"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         'lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /sex="Male"
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Query Match Best Local Similarity

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Score 16; Pred. No.

DB 17; Length 26; 3.3e+04;

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